



PCT

WORLD INTELLECTUAL PROPERTY ORGANIZATION
International Bureau

INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

<p>(51) International Patent Classification ⁶ : C07H 21/04, C12N 15/12, 15/63, 15/64, 15/70, 15/74, 15/90</p>	<p>A1</p>	<p>(11) International Publication Number: WO 96/40722 (43) International Publication Date: 19 December 1996 (19.12.96)</p>
<p>(21) International Application Number: PCT/US96/09006 (22) International Filing Date: 5 June 1996 (05.06.96) (30) Priority Data: 08/482,182 7 June 1995 (07.06.95) US (71) Applicant: CELTRIX PHARMACEUTICALS, INC. [US/US]; 3055 Patrick Henry Drive, Santa Clara, CA 95054-1815 (US). (72) Inventors: MASCARENHAS, Desmond; 27233 Sherlock Road, Los Altos Hills, CA 94022 (US). OLSON, Pamela, S.; 18630 Barnhart Avenue, Cupertino, CA 95014 (US). (74) Agents: PARK, Freddie, K. et al.; Morrison & Foerster L.L.P., 755 Page Mill Road, Palo Alto, CA 94304-1018 (US).</p>	<p>(81) Designated States: AL, AM, AT, AU, AZ, BB, BG, BR, BY, CA, CH, CN, CZ, DE, DK, EE, ES, FI, GB, GE, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, TJ, TM, TR, TT, UA, UG, UZ, VN, ARIPO patent (KE, LS, MW, SD, SZ, UG), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG).</p> <p>Published <i>With international search report.</i> <i>With amended claims.</i></p>	
<p>(54) Title: CHROMOSOMAL EXPRESSION OF HETEROLOGOUS GENES IN BACTERIAL CELLS</p> <p>(57) Abstract</p> <p>The present invention provides compositions and methods for producing a heterologous protein of interest by inserting a copy of a gene encoding the heterologous protein of interest into the chromosome of a host cell, such as <i>E. coli</i>. A chromosomal transfer DNA (a circular, non-self-replicating DNA) is used to integrate the gene encoding the heterologous protein of interest into the host cell chromosome. The chromosomal transfer DNA comprises at least one selectable marker and may optionally include repeated DNA sequences flanking the selectable marker, facilitating chromosomal amplification of the integrated DNA. The gene encoding the protein of interest may be expressed after integration into the chromosome of the host cell; selection for chromosomal amplification may be performed prior to expression of the gene.</p>		

FOR THE PURPOSES OF INFORMATION ONLY

Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

AM	Armenia	GB	United Kingdom	MW	Malawi
AT	Austria	GE	Georgia	MX	Mexico
AU	Australia	GN	Guinea	NE	Niger
BB	Barbados	GR	Greece	NL	Netherlands
BE	Belgium	HU	Hungary	NO	Norway
BF	Burkina Faso	IE	Ireland	NZ	New Zealand
BG	Bulgaria	IT	Italy	PL	Poland
BJ	Benin	JP	Japan	PT	Portugal
BR	Brazil	KE	Kenya	RO	Romania
BY	Belarus	KG	Kyrgyzstan	RU	Russian Federation
CA	Canada	KP	Democratic People's Republic of Korea	SD	Sudan
CF	Central African Republic	KR	Republic of Korea	SE	Sweden
CG	Congo	KZ	Kazakhstan	SG	Singapore
CH	Switzerland	LI	Liechtenstein	SI	Slovenia
CI	Côte d'Ivoire	LK	Sri Lanka	SK	Slovakia
CM	Cameroon	LR	Liberia	SN	Senegal
CN	China	LT	Lithuania	SZ	Swaziland
CS	Czechoslovakia	LU	Luxembourg	TD	Chad
CZ	Czech Republic	LV	Latvia	TG	Togo
DE	Germany	MC	Monaco	TJ	Tajikistan
DK	Denmark	MD	Republic of Moldova	TT	Trinidad and Tobago
EE	Estonia	MG	Madagascar	UA	Ukraine
ES	Spain	ML	Mali	UG	Uganda
FI	Finland	MN	Mongolia	US	United States of America
FR	France	MR	Mauritania	UZ	Uzbekistan
GA	Gabon			VN	Viet Nam

CHROMOSOMAL EXPRESSION OF HETEROLOGOUS
GENES IN BACTERIAL CELLS

TECHNICAL FIELD

5 This invention is related to the field of expression of heterologous genes in bacteria.

BACKGROUND ART

 Genetic engineering has made it possible to produce large amounts of
10 heterologous proteins or polypeptides in bacterial cells by means of recombinant expression systems, especially by expression in such prokaryotes as *Escherichia coli* (*E. coli*) .

 The expressed heterologous proteins may be of mammalian, other eukaryotic, viral, bacterial, cyanobacterial, archaebacterial, or synthetic origin.

15 Unlike native bacterial proteins, which can often be efficiently accumulated within a bacterial cell even when encoded by a single chromosomal gene copy, there are no published reports to date of heterologous proteins being successfully accumulated within bacterial cells to levels exceeding 0.1% of total cell protein when expressed from a single chromosomal gene location.

20 0.1% of total cell protein (150 micrograms protein per trillion bacterial cells) is chosen as a practical measure of successful accumulation of protein because it approximately defines the lower limits of (a) economically significant accumulation of a desired protein by contemporary recombinant bacterial production standards, and (b) visual detection of a protein band by Coomassie-stained polyacrylamide gel analysis of
25 whole bacterial cell extracts.

 The relatively poor performance of non-bacterial genes when expressed in bacterial cells, even when placed under the control of the strongest known bacterial promoters, has been generally attributed to poor translation of the non-bacterial mRNAs

and rapid degradation of newly synthesized non-bacterial proteins. It has almost universally been assumed that, in order to achieve successful accumulation of non-bacterial or heterologous proteins in bacterial cells, the genes encoding the heterologous proteins must be located on multicopy plasmid vectors.

5 A gene carried on one of the multicopy plasmids commonly used for cloning and expressing genes encoding heterologous proteins in *E. coli* usually has a copy number of more than 20 copies/cell. Even low copy number plasmids (e.g., pACYC177 and pLG339) generally exist at 6-10 copies per cell. One disadvantage imposed by plasmid gene dosages is that the expression of even minute amounts of some
10 foreign proteins can kill host cells (see *Meth. Enzymol.* 185:63-65, ed. D. Goeddel, 1990). For this reason, it would be advantageous to reliably limit the copy number of genes encoding such toxic gene products, such as by integrating the gene into the bacterial chromosome at one or a small number of copies per cell. For example, such a system would allow one to make more representative cDNA expression libraries in
15 bacterial hosts if the high-copy expression of one or more of the cDNAs in the library could kill the bacterial host or cause it to grow poorly.

Chromosomal integration of genes encoding heterologous polypeptides would also be advantageous as an alternative means for expression of heterologous proteins in bacterial host cells. Multicopy vectors are often unstable and require the use
20 of antibiotics in the growth medium for maintenance. Present methods of integrating foreign genes into the bacterial chromosome suffer from inefficiency, the inability to control the site of integration of the foreign gene, and/or the inability to control the copy number of the integrated gene. Most importantly, all efforts to date to create recombinant DNA constructs on the bacterial chromosome, wherein a bacterial promoter
25 is fused to a heterologous gene, have involved the creation of viral or plasmid intermediates carrying the construct. Because such intermediates replicate at high copy number, they may be difficult or even impossible to recover in cases where the foreign gene product is toxic to the bacterial cell. Expression of the encoded gene, even at low levels, may be toxic to the host cells, due to the high copy number of these intermediates,
30 which effectively multiplies the level of expression.

Previous methods for achieving the integration of heterologous genes into the chromosome of a bacterial host include the use of phage lambda vectors. The phage DNA in circular form is inserted linearly into the bacterial chromosome by a single site specific recombination between a phage attachment site (attP), 240 bases long, and a
5 bacterial attachment site (attB), only 25 bases long. The two sites have 15 bases in common. This site-specific recombination is catalyzed by a special integrase, specified by the phage gene INT (VIOLOGY pp. 56-57 (Lippincott, 2nd ed., R. Dulbecco and H. Ginsberg, eds., Philadelphia, PA, 1985).

Phage vectors which are INT⁻ can be integrated into the chromosome in a
10 normal fashion as long as integrase is supplied in trans, e.g., by an INT⁺ helper phage (see, e.g., Borck et al. (1976) Molec. Gen. Genet. 146:199-207).

Phage vectors which are both att⁻ and INT⁻ can likewise be integrated into the bacterial chromosome as double lysogens by using att⁺INT⁺ helper phage. Double lysogens are formed by linkage of the prophages at the bacterial attachment site and are
15 integrated into the chromosome by general bacterial recombination between homologous sequences on the defective phage and on the helper phage (see e.g., Struhl et al. (1976) Proc. Natl. Acad. Sci. USA 73:1471-1475). Similarly, it is also possible to integrate non-replicating colE1 replicons into the genome of polA strains of E. coli by means of recombination between the host chromosome and homologous sequences carried by the
20 plasmid vector (Greener and Hill (1980) J. Bacteriol. 144:312-321).

More recently, systems have been specifically designed for the integration of foreign genes into a bacterial host chromosome. For example, U.S. Patent No. 5,395,763 (Weinberg et al.) discloses a chromosomal expression vector for the expression of heterologous genes. This vector was created utilizing a multicopy number
25 plasmid intermediate, into which the gene of interest is cloned, placing the gene in operable linkage with the bacteriophage middle promoter, P_m. This plasmid intermediate, which comprises a defective Mu genome (lacking the genes necessary for the formation of phage particles) is introduced into a packaging strain to produce infectious Mu particles, which are then used to introduce the vector into host cells and
30 integrate the vector into the host cell genome. This vector system is amplifiable once integrated into the host cell genome, but the mechanism of amplification (replicative

transposition) is normally toxic to the host cell, due to integration of the replicating prophage into essential host cell genes (Neidhardt et al., *ESCHERICHIA COLI AND SALMONELLA TYPHIMURIUM: MOLECULAR AND CELLULAR BIOLOGY* (American Society for Microbiology, Neidhardt et al. eds., Washington, D.C., 1987). Because the amplification of this integrated prophage is normally toxic, it is very difficult to obtain and propagate a host cell strain carrying the amplified integrated DNA. This then requires that the gene be amplified each instance that protein production is desired.

Diederich et al. ((1992) "New plasmid vectors for integration into the *attB* attachment site of the *Escherichia coli* chromosome", *Plasmid* 28:14-24) also disclose a system for introducing a gene onto the chromosome of a bacterial host cell. This system utilizes a set of multicopy plasmid vectors which can be integrated into a bacterial chromosome via a phage lambda attachment site. A DNA sequence encoding a promoter operably linked to a gene of interest is cloned into one of the described multicopy number plasmid vectors, the plasmid's origin of replication is removed by restriction enzymes, and the resulting DNA is recircularized and transferred to a host cell, where it integrates into the chromosome.

These new gene transfer systems suffer from the same defect as earlier systems. Both USP 5,395,763 (Weinberg et al.) and Diederich et al. require that the gene of interest be cloned into a multicopy number plasmid while in an operable configuration during the construction of the transfer DNA. The configuration of this multicopy number plasmid makes expression of toxic foreign genes difficult, if not impossible, because the (toxic) gene of interest will be expressed as the multicopy number plasmid is propagated.

Accordingly, there is a need for a method of producing heterologous proteins which can produce large amounts of protein and which minimizes any toxic effect of the heterologous protein to host cells during construction of the producing strain. Applicants have shown surprisingly high protein accumulation (approximately 20% of total cell protein) from expression of low (approximately two) copies of the gene encoding the heterologous protein as shown in Example 2.

SUMMARY OF THE INVENTION

The present invention provides methods and compositions for production of heterologous proteins in bacterial host cells such as E. coli by integrating a chromosomal transfer DNA (a circular, non-self replicating DNA) into the chromosome of a host cell. The chromosomal transfer DNA comprises one or more copies of a gene encoding the heterologous protein of interest.

The present invention, therefore, provides a method for producing a heterologous protein of interest, comprising:

integrating a chromosomal transfer DNA into the chromosome of a host cell such that chromosomal amplification of the integrated DNA is facilitated, the chromosomal transfer DNA comprising at least one copy of a gene encoding a heterologous protein of interest and a selectable marker; and

expressing the gene encoding the heterologous protein of interest, wherein the gene was at no time operably linked to a promoter functional in the host cell in a multicopy number plasmid during the construction of the transfer DNA, and wherein the heterologous protein of interest accumulates to a level of at least 0.1% of total cell protein.

The chromosomal transfer DNA may optionally comprise a promoter operably linked to the gene encoding the heterologous protein of interest, wherein the operable linkage is created by circularization of the chromosomal transfer DNA.

Optionally, the chromosomal transfer DNA may further comprise duplicate DNA flanking the selectable marker. The duplicate DNA may optionally comprise copies of the gene encoding the heterologous protein of interest operably linked to a promoter.

The methods for expression of heterologous proteins may optionally include the step of selecting for chromosomal amplification.

The invention also provides methods for producing a chromosomal transfer DNA, comprising ligating together fragments from a first and a second plasmid vector:

the first plasmid vector comprising a first origin of replication, and a first gene encoding a heterologous protein of interest wherein the first gene is not operably linked to a promoter and a first copy of a duplicate DNA;

the second plasmid vector comprising a second origin of replication, and a first promoter and a second copy of a duplicate DNA;

wherein the origins of replication and the promoter function in the host cell, and wherein either said first plasmid or said second plasmid comprises a selectable marker.

Optionally, the first plasmid may further comprise a second promoter not operably linked to the first gene encoding the heterologous protein of interest and the second plasmid may further comprise a second copy of the gene encoding the heterologous protein of interest not operably linked to the first promoter.

Also provided are chromosomal transfer DNAs for use in production of heterologous proteins of interest, comprising:

a non-bacterial gene of interest operably linked to a promoter functional in a host cell; and

a selectable marker flanked by duplicate DNA,

wherein said gene encoding a heterologous protein is at no time operably linked to a promoter functional in a host cell on a multicopy number plasmid vector during the construction of the transfer DNA.

Optionally, the chromosomal transfer DNA may further comprise two or more copies of the gene encoding the non-bacterial protein of interest, wherein the copies of the gene flank the selectable marker.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 shows steps in the in vitro formation of a chromosomal transfer DNA, a DNA circle which lacks an origin of replication (and thus is incapable of self-replication) and is suitable for integration of a foreign gene into the bacterial chromosome. Until the chromosomal transfer DNA is formed, the foreign gene to be expressed (here an IGF-I fusion gene) is separated from a functional bacterial promoter (here the T7 promoter).

Figure 2 shows a chromosomal transfer DNA formed from the ligation of two DNA fragments. One of the fragments contains a fusion gene comprising sequences encoding *E. coli* DsbA, yeast ubiquitin (beginning with a Met), and human insulin-like growth factor I ("dsbA-ubi-IGF") (not beginning with a Met), as discussed in co-owned, co-pending U.S. patent application no. 08/100,744, filed August 2, 1993. The other DNA fragment contains a T7 promoter. Both the chromosomal transfer DNA and the bacterial chromosome contain a recombination site from phage lambda, att P. The chromosomal transfer DNA is transformed into *E. coli* strain B1384, which makes integrase (INT) under the control of the trp promoter (P-trp). Integrase catalyzes site-specific integration of the chromosomal transfer DNA into the bacterial chromosome at the att site. The trp promoter can be induced during transformation by adding 1 mM indole acrylic acid (IAA) to the medium. Cells with integrated chromosomal transfer DNA sequences are resistant to chloramphenicol (CAM-r, 10 Tg/ml).

Figure 3 shows a B1384 chromosomal integrant resulting from the process described in Figure 2. The integration can be confirmed by amplifying host chromosomal DNA by PCR with various primer sets (e.g., UBUF x IGFR, 1243 x T7REV, or TRPPF x 1239), digesting the amplified fragments with the appropriate restriction enzyme (SacII, HinCII, or BamHI, respectively), and sizing the products by gel electrophoresis).

Figure 4 shows a Western blot of whole cell lysates of chloramphenicol resistant W3110DE3 transductants. Also included are protein size markers (far left lane) and IGF fusion protein (control).

Figure 5 shows a Western blot of whole cell lysates of kanamycin resistant transductants.

Figures 6-9 show diagrammatically the general strategy for construction of chromosomal transfer DNA's. Figure 6 shows a chromosomal transfer DNA comprises a single copy of the gene encoding the heterologous protein of interest and two copies of a second gene which flank the selectable markers, facilitating chromosomal amplification after integration of the chromosomal transfer DNA. Figure 7 shows the "double cassette" system utilized for expression of heterologous proteins in Examples 2 through 6 and Example 8. This embodiment of the chromosomal transfer

DNA comprises two copies of the gene encoding the heterologous protein of interest flanking the selectable markers, facilitating chromosomal amplification of the integrated DNA. Figures 8 and 9 show alternate embodiments of "promoter-less" chromosomal transfer DNAs. These embodiments utilize a DNA sequence homologous to a segment of the host cell chromosome. Integration of promoter-less chromosomal transfer DNAs results in formation of an operably linkage between a host cell promoter and the gene encoding the heterologous protein of interest and the creation of duplicate DNA sequences flanking the selectable markers.

Figures 10-13 show the plasmid genealogy of chromosomal transfer DNAs.

Figure 14 shows the strategy for construction of the two DNA sources used in the double cassette system.

Figures 15 and 16 show the strategy used to construct chromosomal transfer DNAs for integration and expression of the yeast ubiquitin hydrolase gene.

Figure 17 shows the strategy for constructing the chromosomal transfer DNA used to integrate and express a gene encoding a DsbA::ubiquitin::IGF-I fusion protein.

Figure 18 shows the strategy for constructing the chromosomal transfer DNA used to integrate and express a gene encoding a DsbA::2A::IGFBP-3 fusion protein.

Figures 19 and 20 show the strategies used to construct chromosomal transfer DNAs used to integrate and express genes coding for DsbA::2A::IGF-I (Figure 19) and DsbA::3C::IGF-I (Figure 20) fusion proteins.

Figure 21 shows the strategy used to construct the chromosomal transfer DNA used to integrate and express the gene encoding a DsbA::ubiquitin::TGF- β 2 fusion protein.

Figure 22 shows the strategy used to construct the chromosomal transfer DNA used to integrate and express a gene encoding a DsbA::3C::IGFBP-3 fusion protein.

Figure 23 shows an coomassie blue-stained SDS-PAGE gel of whole cell lysates of isolates expressing IGF-I fusion proteins. c49222, c49258#46, and c53063

express a DsbA::ubiquitin::IGF-I fusion protein (left arrow), which is easily visible. Surprising, this high level of expression is seen in c49222 and c49258#46, which were not amplified (i.e. there was no selection for chromosomal amplification of the integrated DNA). c57264#5 and c57264#28 express a DsbA::3C::IGF-I fusion protein while
5 c57265#44 and c57265#54 express a DsbA::2A::IGF-I fusion protein. Again, the expressed fusion protein is easily visible. Densitometric analysis of this gel indicates that all of the isolates accumulate protein in excess 19% of total cell protein (average protein accumulation is 25.7% of total cell protein).

Figure 24 shows a Southern blot of chromosomal DNA isolated from
10 c49222, c49258#46, c53063, c57264#5, c57264#28, c57265#44, and c57265#54. The blot was probed with a DNA fragment encoding ubiquitin fused to IGF-I. The higher molecular weight band in each lane represents a single copy of the integrated IGF-I fusion protein gene in each isolate. The lower molecular weight band also represents the integrated IGF-I fusion protein gene, but this fragment can be amplified by chromosomal
15 amplification. Isolates c53063, c57264#5, c57264#28, c57265#44, and c57265#54 have clearly been amplified, showing about 3 to 5 fold amplification.

Figure 25 shows coomassie blue-stained SDS-PAGE gels showing protein accumulation in isolates carrying integrated genes encoding IGFBP-3 fusion proteins. A) shows protein accumulation in an isolate expressing a DsbA::2A::IGFBP-3 fusion
20 protein. The right lane shows protein expression after induction of T7 RNA polymerase by addition of IPTG to the culture medium. B) shows protein accumulation in an isolate expressing a DsbA::3C::IGFBP-3 fusion protein. As in Figure 23, the bands representing the fusion protein are easily visible. Densitometric scanning of these gels found that the accumulated protein represented 22.6% in Panel A, and the two isolates in Panel B
25 accumulated 33.% and 28.2% of total cell protein (left to right, respectively).

Figure 26 shows a coomassie blue-stained SDS-PAGE gel showing protein accumulation from host cells expressing a gene encoding
DsbA::ubiquitin::TGF- β 2. M indicates molecular weight markers and C indicates a
positive control. The two Plasmid lanes (Lanes 1 and 2) are used as a standard to
30 compare protein accumulation from multicopy number plasmid vectors to protein accumulation from genes integrated into the chromosome. Lanes 3 and 4 are whole cell

lysates of isolates which were negative for T7 RNA polymerase activity when streaked against phage 4107. Densitometric analysis of this gel showed that the plasmid strain accumulated protein to 26.4% of total cell protein. Protein accumulation was measured for isolates 48, 56, 59, 65 and 66, and showed protein accumulation to 36.7%, 33.3%,
5 32.1%, 29.5%, and 26.7%, respectively.

MODES FOR CARRYING OUT THE INVENTION

The present invention resides in (a) the creation of an operable linkage between a promoter and a gene encoding a heterologous protein of interest with the
10 linkage being formed either during the construction of a chromosomal transfer DNA or as a result of its integration into the host cell chromosome and (b) the simultaneous creation of a means for the appropriate chromosomal amplification of the integrated gene of interest.

In the preferred embodiments, the creation of the chromosomal transfer
15 DNA simultaneously achieves two goals; (1) the operable linkage of the promoter and the gene of interest and (2) the positioning of duplicate DNA sequences flanking a selectable marker (which can function as a means to facilitate the amplification of the chromosomal transfer DNA). Another embodiment creates the operable linkage between the gene and the promoter during creation of the chromosomal transfer DNA, while the
20 means for chromosomal amplification (duplicate DNA sequences flanking the chromosomal transfer DNA) is created as a result of the integration.

Other methods can achieve either or both of these results by integration of a chromosomal transfer DNA into a suitable site on the chromosome. For example, integration of a gene of interest near a promoter on the bacterial chromosome can be
25 designed to result in an operable linkage (for example, by integrating a chromosomal transfer DNA into an operon on the host cell chromosome). The site of integration or sequences adjacent to the site of integration may facilitate amplification (e.g. where the site is located in a transposable element, by providing duplicate DNA sequences, or even by providing a region of DNA sequence homologous to a portion of the chromosomal
30 transfer DNA, thus providing duplicate DNA sequences).

The present invention employs "chromosomal transfer DNA" which may be used to simply, efficiently, and reliably insert a copy of a heterologous gene into the chromosome of a host cell, e.g., *E. coli*. A chromosomal transfer DNA is a circular DNA comprising one or more copies of a gene encoding a heterologous protein of interest, a selectable marker (e.g., an antibiotic resistance gene), a recombination site (e.g., a site-specific recombination site such as lambda attP or attB or a DNA sequence homologous to a segment on the host cell chromosome), and means for facilitating the amplification of the chromosomal transfer DNA following recombination into the host cell chromosome, and lacking an origin of replication or autonomously replicating sequence (ARS). The chromosomal transfer DNA is therefore incapable of replicating independently when introduced in to the host cell. The chromosomal transfer DNA may optionally carry a promoter operably linked to the gene of interest.

When a chromosomal transfer DNA carrying a site-specific recombination site is introduced into a host cell having a chromosome which contains a second, similar recombination site (e.g., another attP or attB site), expression in the host cell of an enzyme which is capable of catalyzing the site-specific recombination of the recombination sites (e.g., integrase) results in the integration of the vector into the host cell chromosome at the recombination site. This site-specific recombination process is much more efficient than general recombination systems acting on homologous vector and host chromosomal sequences and results in integrated sequences having greater stability, particularly when integrase synthesis can be controlled. Integrase may also be provided by a plasmid or other DNA molecule transiently or stably present in the host cell at the time when the chromosomal transfer DNA is introduced.

It will be apparent to one skilled in the art that there are a variety of methods other than the preferred method utilizing attP, attB, and INT which may be used to integrate a chromosomal transfer DNA into the chromosome of a host cell. For example, non-replicating colE1 replicons, transposable elements, or even naked DNA carrying sequences homologous to sequences found on the host chromosome may be used to insert the chromosomal transfer DNA into the host chromosome. The multicopy colicin plasmids ColE1, CloDF13, ColK, and ColA all comprise site-specific recombination systems including a cis- and trans-acting element. For use in the present

invention, the cis-acting element from one of these plasmids may be included on the chromosomal transfer DNA and the trans-acting element may be on the chromosomal transfer DNA or provided by the host cell. Transposons, such as the insertion sequence (IS) and Tn3 families of transposons may be used to integrate DNA into the chromosome of a host cell. As with the colicin plasmids described above, the cis-acting transposon elements are included on the chromosomal transfer DNA, while the trans-acting factor may be included on the chromosomal transfer DNA or provided by the host cell. The chromosomal transfer DNA may also carry a DNA sequence homologous to a sequence found on the host cell chromosome, facilitating integration of the chromosomal transfer DNA by homologous recombination. All of these methods fall within the scope of the invention.

An important feature of this approach is that the gene encoding the heterologous protein of interest is at no time operably linked to a functional promoter on a multicopy vector during construction of the transfer DNA. By keeping a functional promoter separated from the gene of interest until immediately before the foreign gene is introduced into the cell at low copy number, the potential toxic or lethal effects of the gene product can be minimized. A toxic foreign gene will not be expressed from a multicopy number plasmid if the gene is not operably linked to a promoter. Other methods for integrating a gene of interest into the host cell chromosome utilize multicopy number plasmids carrying a gene of interest operably linked to a promoter (e.g., Diederich et al. and Weinberg et al.); these genes will be expressed during the propagation of the plasmid, making it extremely difficult, if not impossible, to produce sufficient quantities of the plasmid if the gene of interest is toxic to the host cells in which the plasmid is propagated.

The operable linkage between the gene encoding the heterologous protein of interest and the promoter may be created as a result of the formation of the chromosomal transfer DNA or as a result of integration into the host cell chromosome. In the case where the operable linkage is formed as a result of the formation of the chromosomal transfer DNA, the linkage is created by circularization of the chromosomal transfer DNA. Circularization may be accomplished by, for example, ligation of one or more DNA fragments to form a circular DNA or by homologous recombination into a

circular DNA, which would result in circularization of the insert. Preferably, circularization is accomplished by ligation of one or more DNA fragments.

Alternatively, high level expression of less toxic gene products can be accomplished by multiple integrations or by selection for amplification of integrated genes.

Recombinant DNA Methods and Reagents

General techniques for nucleic acid manipulation useful for the practice of the claimed invention are described generally, for example, in Sambrook et al., MOLECULAR CLONING: A LABORATORY MANUAL, Vols. 1-3 (Cold Spring Harbor Laboratory Press, 2 ed., (1989); or F. Ausubel et al., CURRENT PROTOCOLS IN MOLECULAR BIOLOGY (Green Publishing and Wiley-Interscience: New York, 1987) and periodic updates. Reagents useful in nucleic acid manipulation, such as restriction enzymes, T7 RNA polymerase, DNA ligases and so on are commercially available from such vendors as New England BioLabs, Boehringer Mannheim, Amersham, Promega Biotec, U.S. Biochemicals, and New England Nuclear.

Definitions

"Foreign" or "heterologous" or "non-bacterial:" "native" or "homologous" A "foreign or "heterologous" polypeptide is a polypeptide which is not normally found in a host cell of a particular species. The nucleic acid encoding such a polypeptide is also referred to as "foreign" or "heterologous." For example, insulin-like growth factor (IGF), insulin-like growth factor binding protein (IGFBP), and transforming growth factor-beta (TGF- β) are native to mammalian cells and human rhinovirus 3C protease is native to viruses and virally-infected mammalian cells, but these proteins are foreign or heterologous to *E. coli*. A "non-bacterial protein" is a protein or polypeptide which is not naturally found in a bacterial cell. Non-bacterial proteins include viral and eukaryotic proteins. Non-bacterial, foreign, or heterologous proteins may also be fusions between non-bacterial, foreign, or heterologous proteins and other proteins or polypeptides. For the embodiments encompassed by this invention, both "heterologous protein" and "non-bacterial protein" may be expressed. As disclosed

herein, genes encoding heterologous or non-bacterial proteins of interest do not contain promoters functional in the host cell. The genes must be linked to a separate promoter that is functional in the host cell in order to be expressed. A "native" or "homologous" polypeptide or DNA sequence, by contrast, is commonly found in the host cell. A promoter or other sequence effecting, for example, the transcription or translation of a gene is also considered "homologous" if it is functional in the host cell. For example, a T7 promoter is considered "homologous" to an *E. coli* host cell, since, if T7 RNA polymerase is present in the cell, the T7 promoter is capable of driving the transcription of a polypeptide-encoding sequence to which it is operably linked.

10 "Genes encoding heterologous, foreign or non-bacterial proteins" "Genes encoding heterologous, foreign or non-bacterial proteins" contain all of the genetic elements necessary for the expression of the heterologous, foreign or non-bacterial protein with the exception of a promoter functional in the host cell. These genes encompass recombinant genes which may include genetic elements native to the host cell. Further, the coding regions of these genes may optionally be optimized for the codon usage of the host cell.

"Encode" A nucleic acid is said to "encode" a polypeptide if, in its native state or when manipulated by recombinant DNA methods, it can be transcribed and/or translated to produce the polypeptide.

20 "Operably linked" A nucleic acid sequence is operably linked when it is placed into a functional relationship with another nucleic acid sequence. For example, a promoter is operably linked to a coding sequence if the promoter affects its transcription or expression. Generally, DNA sequences which are operably linked are contiguous and, where necessary, in reading frame.

25 "Recombinant" A "recombinant" nucleic acid is one which is made by the joining of two otherwise separated segments of nucleic acid sequence *in vitro* or by chemical synthesis.

"Chromosomal amplification" "Chromosomal amplification" refers to the increase in copy number of a DNA sequence on the host chromosome. Chromosomal amplification does not refer to extrachromosomal amplification such as replication of

30

multicopy number plasmids or in vitro amplification such as the polymerase chain reaction (PCR).

Probes and primers

5 Nucleic acid probes and primers are isolated nucleic acids, generally single stranded, and, especially in the case of probes, are typically attached to a label or reporter molecule. Probes are used, for example, to identify the presence of a hybridizing nucleic acid sequence in a tissue or other sample or a cDNA or genomic clone in a library. Primers are used, for example, for amplification of nucleic acid
10 sequences, e.g., by the polymerase chain reaction (PCR). The preparation and use of probes and primers is described, e.g., in Sambrook et al., supra or Ausubel et al. supra.

Chemical synthesis of nucleic acids

 Nucleic acids, especially short nucleic acids such as amplification
15 primers, may be produced by chemical synthesis, e.g., by the phosphoramidite method described by Beaucage and Carruthers (1981) Tetra. Letts. 22:1859-1862 or the triester method according to Matteucci et al. (1981) J. Amer. Chem. Soc. 103:3185, and may be performed on automated oligonucleotide synthesizers. A double-stranded fragment may be obtained from the single-stranded product of chemical synthesis either by synthesizing
20 the complementary strand and annealing the strands together under appropriate conditions or by adding the complementary strand using DNA polymerase with an appropriate primer sequence.

Features of chromosomal transfer DNA and of plasmids used in their construction

25 Chromosomal transfer DNA comprises a DNA fragment encoding a selectable marker and a sequence encoding a desired heterologous polypeptide. Optionally, a chromosomal transfer DNA may also comprise, in an operable linkage to the sequence encoding the desired heterologous polypeptide, transcription and translation initiation regulatory sequences and expression control sequences, which may include a
30 promoter, an enhancer and necessary processing information sites, such as ribosome-binding sites, and mRNA stabilizing sequences, as well as any necessary

secretion signals, where appropriate, which allow the protein to cross and/or lodge in cell membranes, and thus attain its functional topology, or be secreted from the cell.

Plasmids used in construction of a chromosomal transfer DNA will also typically comprise a replication system recognized by the host, including an origin of replication or autonomously replicating sequence (ARS). In the case where a plasmid used in the construction of a chromosomal transfer DNA carries duplicate DNA sequences, the plasmid may be propagated in a rec⁻ host cell. Preferably, rec⁻ host cells are used for propagation of plasmids used to create chromosomal transfer DNAs and plasmids carrying components of chromosomal transfer DNAs when these plasmids carry duplicate DNA sequences, and are not generally utilized as host cells for integration of chromosomal transfer DNAs.

Chromosomal transfer DNA may be prepared from such vectors by means of standard recombinant techniques well known in the art and discussed, for example, in Sambrook et al., supra or Ausubel et al. supra.

An appropriate promoter and other sequences necessary for efficient transcription and/or translation are selected so as to be functional in the host cell. Examples of workable combinations of cell lines and expression vectors are described in Sambrook et al., supra or Ausubel et al., supra; see also, e.g., Metzger et al. (1988) Nature 334:31-36. Promoters such as the trp, lac and phage promoters (e.g., T7, T3, SP6), tRNA promoters and glycolytic enzyme promoters are useful in prokaryotic hosts. Useful yeast promoters include the promoter regions for metallothionein, 3-phosphoglycerate kinase or other glycolytic enzymes such as enolase or glyceraldehyde-3-phosphate dehydrogenase, enzymes responsible for maltose and galactose utilization, and other. See, e.g., Hitzeman et al. EP 73,657A. Appropriate mammalian promoters include the early and late promoters from SV40 (Fiers et al. (1978) Nature 273:113) or promoters derived from murine Moloney leukemia virus, mouse mammary tumor virus, avian sarcoma viruses, adenovirus II, bovine papilloma virus or polyoma virus. In addition, the construct may be joined to an amplifiable gene (e.g., DHFR) so that multiple copies of the gene may be made, where desired. For appropriate eukaryotic enhancer and other expression control sequences see, e.g.,

ENHANCERS AND EUKARYOTIC GENE EXPRESSION (Cold Spring Harbor Press, New York, 1983).

It is preferable that the promoter driving expression of the heterologous gene when integrated in the chromosome of the host is controllable.

5 Chromosomal transfer DNAs and plasmids employed in their construction generally comprise a selectable marker, a gene encoding a protein necessary for the survival or growth of a host cell transformed with the chromosomal transfer DNA or plasmid. Typical selectable markers (a) confer resistance to antibiotics or other toxic substances, e.g. ampicillin, neomycin, methotrexate, etc.; (b) complement auxotrophic
10 deficiencies; or (c) supply critical nutrients not available from complex media, e.g. the gene encoding D-alanine racemase for Bacilli. The choice of the proper selectable marker will depend on the host cell.

The chromosome transfer DNAs of the present invention may contain a site-specific recombination site, such as the phage lambda attP site. When transformed
15 into a bacterial host strain (such as E. coli B1384) which makes the enzyme integrase, integrase recognizes the attP site on the chromosomal transfer DNA and catalyses its recombination with an att site (integrase can catalyze a recombination between two attP and attB or two attP sites). Bacterial host cells bearing the integrated DNA are selected for on the basis of a selectable marker carried on the integrated DNA.

20 Thus, integration utilizing site-specific recombination generally involves expression of an enzyme such as integrase which can catalyze site-specific recombination and the presence of a site recognized by the enzyme on both the chromosomal transfer DNA and the bacterial chromosome. Other site-specific recombination systems characterized by an "integrase" or similar enzyme and sites specifically recognized by
25 the "integrase" could be used as well.

High level expression of a foreign gene integrated into the chromosome of a host cell in multiple copies is also possible, e.g., by incorporating multiple att sites in the host cell chromosome and introducing multiple chromosomal transfer DNAs into the host cell. Additionally or alternatively, host cells containing multiple copies of the
30 integrated DNA may be obtained by selecting for chromosomal amplification. Chromosomal amplification is facilitated when the selectable marker is flanked by

duplicate DNA sequences. Preferably, the duplicate DNA sequences flank a first and a second selectable marker. The first selectable marker is effective at low copy number and can be used to select for integration of the chromosomal transfer DNA. The second selectable marker is preferably effective only at high copy number. Following selection
5 for integration using the first selectable marker, the second selectable marker is then used to select for host cells which contain multiple copies of the integrated DNA.

An important feature of the chromosomal transfer system of the present invention is that the gene encoding the heterologous protein is not expressed before integration; it is not operably linked to a promoter until either (a) the transfer DNA is
10 constructed in vitro or (b) the chromosomal transfer DNA is integrated into the host cell chromosome. This approach allows one to employ high copy number plasmids as DNA sources in constructing the chromosomal transfer DNA. High copy number plasmids carrying a toxic heterologous gene are often difficult to propagate when the toxic gene is operably linked to a promoter. Low copy number plasmids are more difficult to work
15 with in the laboratory. For example, DNA minipreps may produce inadequate DNA for in vitro manipulations. The chromosomal transfer DNA is constructed from one or more DNA sources by circularization of selected DNA fragments.

When a single DNA is used to construct the chromosomal transfer DNA, both the gene encoding the heterologous protein of interest and the promoter are located
20 on the same DNA, however the gene and promoter are not operably linked. This may be accomplished by, for example, placing the promoter and gene of interest on either side of a spacer DNA sequence which blocks any operable linkage (for example, by including a terminator sequence). Preferably, this intervening DNA sequence also includes any other portions of the source DNA which must be removed for creation of the chromosomal
25 transfer DNA, such as an origin of replication or ARS. The chromosomal transfer DNA is constructed by deleting the DNA sequence which blocks the operable linkage between the gene and the promoter, then circularizing the remaining DNA.

As shown in Figures 7, 8, and 9, chromosomal transfer DNAs may optionally include a DNA sequence for the expression of *E. coli* cyclophilin, as described
30 in U.S. Patent No. 5,459,051.

There are several methods by which one may construct a chromosomal transfer DNA using two or more DNA sources. In one preferred embodiment, shown in Figure 7, also uses two DNA sources. In this embodiment, each of the two DNA sources carries a copy of the gene encoding the heterologous protein of interest and the promoter, but the gene encoding the heterologous protein of interest and promoter are not operably linked on either DNA source. As with the previously described embodiment, other necessary sequences may be carried by either DNA source (alternatively the other necessary sequences may be provided by one or more accessory DNA sources). The two DNA sources are cleaved, then joined to each other, forming a circular chromosomal transfer DNA which has two copies of the foreign gene, each operably linked to a copy of a promoter. The promoter from the first DNA source is operably linked to the gene encoding the heterologous protein of interest from the second DNA source, and the promoter from the second DNA source is operably linked to the gene encoding the heterologous protein of interest from the first DNA source.

Chromosomal transfer DNAs may also be designed without promoters (Figures 8 and 9). These promoter-less chromosomal transfer DNAs are integrated into target sites on the bacterial chromosome which place the gene encoding the heterologous protein of interest into an operable linkage with a promoter on the host cell chromosome. The chromosomal transfer DNA of this embodiment includes a copy of a gene encoding a heterologous protein of interest linked in-frame to a segment of target-site DNA segment homologous to DNA on the host cell chromosome and a selectable marker. This target site DNA sequence will typically be the 5' end of a gene located on the bacterial chromosome downstream from a promoter. Integration of the chromosomal transfer DNA into the host cell chromosome will place the gene encoding the heterologous protein of interest into operable linkage with a bacterial promoter. The target sequence on the host cell chromosome may be a naturally occurring sequence or may be a site which is introduced into the chromosome of the host cell. A target may be introduced into the chromosome of a host cell utilizing a DNA sequence homologous to a segment of the host cell chromosome, as described above for integration of the chromosomal transfer DNA. A target site may also be introduced using site-specific recombination, such as the attB/attP/INT system described above. A target site sequence

is at least about 10 bases long, preferably at least about 30 bases long, and most preferably at least about 100 bases long. The DNA sequence on the chromosomal transfer DNA and the target site are at least about 80% homologous, preferably at least about 90% homologous and most preferably at least about 95% homologous. A target site is preferably rare in the host cell chromosome and, more preferably, is unique in the host cell chromosome. Integration of the chromosomal transfer DNA using a sequence homologous to a segment on the host cell chromosome facilitates amplification of the integrated DNA by placing duplicate DNA sequences flanking the integrated DNA (see Figures 8 and 9).

10 Introducing DNA into host cells

A variety of methods for introducing nucleic acids into host cells are known in the art, including, but not limited to, electroporation; transfection employing calcium chloride, rubidium chloride calcium phosphate; DEAE-dextran, or other substances; microprojectile bombardment; lipofection; and infection (where the vector is an infectious agent, such as a retroviral genome). See generally, Sambrook et al., supra and Ausubel et al., supra.

Host cells

The methods of the present invention are preferably used with prokaryotic host cells, although they would be applicable to eukaryotic host cells as well. Among prokaryotic hosts, gram negative bacteria are preferred, especially Escherichia coli. Other prokaryotes, such as Bacillus subtilis or Pseudomonas may also be used.

Mammalian or other eukaryotic host cells, such as yeast, filamentous fungi, plant, insect, amphibian or avian species may also be used. See, TISSUE CULTURE (Kruse and Patterson, ed., Academic Press, 1973). Useful mammalian host cell lines include, but are not limited to, VERO and HeLa cells, Chinese hamster ovary (CHO) cells, and W138, BHK, and COS cell lines.

Amplification of Integrated DNA

30 Amplification of integrated genes can be efficiently accomplished by any of several methods, for example, chromosomal duplication or replicative transposition.

- Integrated DNA which contains or is flanked by duplicate DNA sequences of 25 or more base pairs will form chromosomal duplications (Normark et al. (1977) J. Bacteriol. 132:912-922; Edland et al. (1979) Mol. Gen. Genet. 173:115-125; Tlsty et al. (1984) Cell 37:217-224; Stern et al. (1984) Cell 37:1015-1026). Selection for duplications
- 5 (amplification) is greatly facilitated if the duplicate DNA contains a selectable marker, such as an antibiotic resistance gene or a gene which complements a host cell deficiency. Preferably the integrated DNA includes two selectable markers; a first selectable marker which is operable at low copy number and is used to select for integrants, and an second selectable marker which requires high copy number and is used to select for host cells
- 10 which have amplified the integrated DNA. Amplification may also be accomplished by replicative transposition, in the case where the chromosomal transfer DNA contains the appropriate transposon sequences or the chromosomal transfer DNA is integrated into a transposon. Preferably, amplification is accomplished by selection for chromosomal duplications.

Production of non-bacterial proteins

Following integration of the chromosomal transfer DNA into the host cell chromosome, and optionally following amplification of the integrated DNA, the foreign gene may be expressed, resulting in the production of the non-bacterial protein of interest. It is preferable that the promoter controlling expression of the integrated gene be controllable (i.e., inducible), so that any toxic effects of the gene product can be minimized. Following expression of the foreign gene, the protein product may be purified. As will be apparent to one skilled in the art, the purification method used will depend on the identity of the foreign protein.

The invention will be better understood by reference to the following examples, which are intended to merely illustrate the invention. The scope of the invention is not to be considered limited thereto.

EXAMPLES

Example 1

Integration of a chromosomal transfer DNA comprising a foreign gene into the chromosome of *E. coli* strain B1384

The general strategy for integrating a chromosomal transfer DNA comprising a foreign gene into the chromosome of *E. coli* is depicted schematically in Figure 1. Two plasmids were constructed: pDM25432 contained a foreign gene of interest (in this example, an IGF-I fusion gene) lacking an operably linked bacterial promoter; pDM25423 contained a T7 promoter. By ligating restriction fragments purified from each of these vectors, a DNA circle lacking an origin of replication - a chromosomal transfer DNA - was generated. This chromosomal transfer DNA contained an antibiotic resistance gene which affords resistance to chloramphenicol (CAM-r) and a site-specific recombination site from phage lambda, *attP*. This chromosomal transfer DNA is transformed into a bacterial strain such as *E. coli* B1384 (Mascarenhas et al. (1983) *Virology* 124:100-108) (Figure 2), which makes the enzyme integrase (INT) under the control of the *trp* promoter, which can be induced during transformation by adding 1 mM indole acrylic acid (IAA) to the medium. B1384 also contains an *attP* in its chromosome. Integrase recognizes the *attP* sites on the

chromosomal transfer DNA and in the chromosome of B1384 and catalyses their recombination, leading to the site-specific integration of the chromosomal transfer DNA into the bacterial chromosome at the att P site (Weisberg et al. Comprehensive Virology, vol. 8, pp. 197-258 (Plenum, Fraenckel-Conrat and Wagner, eds., New York, NY, 1977).

- 5 Bacterial host cells bearing the integrated DNA are selected for on the basis of their resistance to chloramphenicol.

Chloramphenicol-resistant chromosomal integrants were tested as summarized in Figure 3. The presence of the integrated chromosomal transfer DNA was confirmed by amplifying host chromosomal DNA by PCR with the following primer sets

- 10 (e.g., UBUF x IGFR, 1243 x T7REV, or TRPPF x 1239)

IGFR: 5' ... CCC ATC GAT GCA TTA AGC GGA TTT AGC
CGG TTT CAG...3'

- 15 #1239: 5' ...GCC TGA CTG CGT TAG CAA TTT AAC TGT
GAT...3'

#1243: 5' ...CTG GGC TGC TTC CTA ATG CAG GAG TCG
CAT...3'

- 20 #1227: 5' ...TAA TAC GAC TCA CTA TAG GGA GA...3'

TRPPF: 5' ...GAT CTG TTG ACA ATT AAT CAT CGA ACT
AGT TAA CTA GTA CGC AAG TT...3'

- 25 T7REV: 5' ...TGC TAG TTA TTG CTC AGC GG...3'

CYCF1: 5' ...CAG GAT CCG ATC GTG GAG GAT GAT TAA
ATG GCG AAA GGG GAC CCG CAC...3'

- 30 CYCR1: 5' ...CAG GAA GCT TAC GGC AGG ACT TTA GCG
GAA AG...3'

UBUF: 5' ...GGG GCC GCG GTG GCA TGC AGA TTT TCG
TCA AGA CTT TGA...3'

35

The amplified fragments were digested with the appropriate restriction enzyme (SacII, HincII, or BamHI, respectively). The products were sized by agarose gel electrophoresis. Presence of the integrated sequences was demonstrated by amplification of:

chromosomal ubiquitin and IGF sequences, demonstrating the presence of the relevant foreign gene;

- chromosomal tet and T7 sequences, demonstrating the juxtaposition of the T7 promoter and the fusion gene; and
- 5 ● adjacent chromosomal trp and tet sequences, demonstrating insertion of the chromosomal transfer DNA at the expected location.

The chromosomal integration of the chromosomal transfer DNA was also confirmed by the following evidence:

- resistance of the bacterial host to chloramphenicol;
- 10 ● no plasmid DNA in DNA minipreps;
- lack of beta-lactamase enzymatic activity, confirming the absence of the parental plasmids (beta-lactamase was assayed using a chromogenic substrate, 7-thienyl-2-acetamido-3-2-4 n,n-dimethylaminophenylazopyridiniummethyl-3cephem-4 carboxylic acid (PADAC), as described in ENZYME INHIBITORS pp. 169-177 (Verlage Chemie, Broderick, V., ed.); and
- 15 ● segregation analysis: Isolates were grown in L broth with or without 1 mM IAA at 37° C overnight and plated on LB agar plates. Single colonies from each culture were tested for retention of
- 20 chloramphenicol resistance. 100% retention was observed from cultures without IAA; 11% retention was observed in cultures with IAA.

Six of seven isolates tested showed the expected phenotypes.

- B1384 does not contain the gene for T7 RNA polymerase. In order to test the expression of the chromosomal constructs, P1 lysates were prepared on each of the
- 25 six strains carrying the integrated DNA and used to transduce strain W3110DE3 to chloramphenicol resistance (A SHORT COURSE IN BACTERIAL GENETICS: A LABORATORY MANUAL AND HANDBOOK FOR ESCHERICHIA COLI AND RELATED BACTERIA (Cold Spring Harbor Laboratory Press, Miller, J.H., ed., 1992)). Strain W3110DE3 carries the T7 RNA polymerase gene under the control of the lac promoter. It is also Gal⁺, unlike
- 30 B1384. Transductants were therefore selected on galactose minimal plates containing 20 Tg/ml chloramphenicol. Single colonies from each transduction experiment (independent donors) were purified and tested further.

The results obtained were identical in all six independent cases: the chromosomal transfer DNA was transferred with high efficiency to a new location on the bacterial chromosome, the att sites flanking the prophage in W3110DE3. This was confirmed by

- 5 ● chloramphenicol resistance;
- no plasmid DNA in DNA minipreps;
- i21 immunity (DE3 lysogen; phage lysates were plated on bacterial lawns by standard techniques);
- gal⁺ (i.e. growth on galactose minimal plates);
- 10 ● expression of IGF protein under lac control (expression and analysis carried out as described in Example 1 or co-owned, co-pending U.S. patent application Serial No. 08/101,506, filed August 2, 1993).

Chromosomal DNA from the six strains ("integrants") was digested to completion with BglII and NcoI and a Southern blot of the digested DNA was probed with a labeled 0.6 kb DsbA DNA probe which covers the entire gene sequence coding for mature DsbA (Bardwell et al. (1991) Cell 67:581-589; see also Kamitani et al. (1992) EMBO J 11:57-62). Each of the six integrants contained insertions; the blots demonstrated the existence of several double insertions, one single insertion, and one (isolate WB3-6) apparently duplicate double (i.e. triple) insertion.

The six integrants were tested for expression of the IGF fusion protein after induction with isopropyl-J-thiogalactopyranoside (IPTG). Cells were induced with IPTG for two hours and whole cell extracts for the induced integrants, as well as size markers and an IGF fusion protein control, were separated by 12% SDS-PAGE, Western blotted, and reacted with polyclonal anti-IGF sera (see Example 1 of co-owned, co-pending U.S. patent application Serial No. 08/101,506, filed August 2, 1993) (Figure 4). Isolate WB3-6 (Figure 4, lane 6) showed the highest levels of expression of the IGF fusion protein. An induced band of the same size was also seen on Coomassie blue-stained gels.

30 A different binary system was used to generate a chromosomal transfer DNA carrying a kanamycin resistance marker. The plasmids used, pDM25424 and pDM25427, are described in the figures. The configuration and location of the insert

were confirmed by PCR, giving results which were virtually identical to those described above. After transduction into the W3110DE3 background, several individual isolates were obtained which expressed the IGF fusion protein at level that could be easily detected by Western blotting (Figure 5). Procedures used were identical to the ones
5 described above for the chloramphenicol-resistant isolates, except that the antibiotic and resistance gene were kanamycin instead of chloramphenicol. Purified fusion protein was the control. Lanes 1 and 2 contain whole cell lysates from two transduced isolates.

The construction of the vectors employed in the two binary systems is summarized in Figures 10-13. The sources for the plasmids employed were: pBR322,
10 pUC18, pUC19, pKK233-2, pRC99A, pCH110, and pNEO (Pharmacia, Piscataway, NJ); pLG339HLY (Dr. Barry Holland, Institute de Génétiques et Microbiologie, Université Paris-Sud); pRcCMV (Invitrogen, San Diego, CA); pACYC177 and pACYC184 (New England BioLabs, Beverly, MA); pET3b (Studier and Moffat (1986) *J. Mol. Biol.* 189:113-130); pYZ22070 (described in Example 1 of co-owned, co-pending
15 U.S. patent application Serial No. 08/101,744, filed August 2, 1993).

E. coli K-12 strain W3110 was obtained from B. Bachmann, ECGSC, Yale University. It was lysogenized with the DE3 defective phage as described by Studier and Moffat (1986) *J. Mol. Biol.* 198:113-130. W3110DE3 was one such lysogen. The cyclophilin gene was amplified by the polymerase chain reaction (PCR)
20 from W3110 using the primers CYCF1 and CYCR1 (see above).

Example 2

Chromosomal expression of a DsbA::ubiquitin::IGF-I fusion gene

A DsbA::ubiquitin::IGF-I fusion gene was assembled and integrated into
25 the chromosome of bacterial host cells with a chromosomal transfer DNA produced using the double-cassette binary system. The strategy for constructing the double cassette binary system vectors is shown in Figure 14. The general strategy for constructing a chromosomal transfer DNA (CTD) with the double cassette system is shown in Figure 7. The strategy used to create the chromosomal transfer DNA carrying
30 the DsbA::ubiquitin::IGF-I fusion gene is shown in Figure 12. Following chromosomal integration, the fusion gene was expressed, resulting in extremely high levels of protein accumulation.

The double cassette binary system utilizes two plasmids, pDM25470 and pDM25465, as shown in Figures 7 and 14. pDM25425 is a pUC19 derivative carrying a copies of attP, the T7 promoter, and a copy of the rrn1t2 terminator, from which a 1.6 kb fragment was deleted by BglII/BamHI digestion. A terminator and a sequence encoding DsbA (a 1.5 kb NcoI(fill)/NsiI fragment from pDM25463) was added ligated to EcoRI(fill)/NsiI-digested pDM25459 to form pDM25470 (one of the double cassette binaries). The other double cassette plasmid, pDM25465, carries two copies of a terminator, a kanamycin resistance gene, and the cyclophilin gene (the use of the cyclophilin gene to aid in protein production is described in co-owned, co-pending U.S. patent application Serial Number 08/101,506, incorporated herein by reference in its entirety). The cyclophilin gene was cloned from pER15951 (HindIII(fill)/XbaI, 0.6 kb fragment) into pDM25424 (BamHI(fill)/XbaI, 5.2 kb fragment; a pUC19 backbone carrying two copies of a terminator and a kanamycin resistance gene). The kanamycin resistance gene in pDM25430 (derived from pDM25424) was insufficiently effective, so it was replaced with a kanamycin resistance gene from pLG339hly (PvuII/EcoRI digest), creating plasmid pDM25443. The T7 promoter was cloned into pDM25443 by annealing oligos T7F and T7R and ligating them the EcoRI-digested pDM25443, creating pDM25465.

Two sets of oligonucleotides were synthesized (1, 2, 1R, 2R and 3, 4, 3R, 4R), phosphorylated, denatured, and annealed. The annealing product of 1, 2, 1R, and 2R, which encodes ubiquitin, was ligated into pUC18 (SphI-BamHI digest). The annealing product of 3, 4, 3R, and 4R, which encodes IGF-I, was ligated into pUC18 (EcoRI-BamHI digest). The resulting plasmids were transformed into JM109 and the transformed host cells were selected on ampicillin plates. Transformants were analyzed for the presence of the ubiquitin and IGF-I sequences, then sequenced to identify correctly formed constructs. One isolate from each was selected, and designated pPO39354 and pPO39334, respectively.

1
30 5'-CAG ATT TTC GTC AAG ACT TTG ACC GGT AAA ACC ATA
ACA TTG GAA GTT GAA CCT TCC GAT ACC ATC GAG AAC GTT
AAG GCG AAA ATT CAA GAC AAG GAA GGT ATC CCT CCA
GAT CA-3'

2
5'-ACA AAG ATT GAT CTT TGC CGG CAA GCA GCT AGA AGA
CGG TAG AAC GCT GTC TGA TTA CAA CAT TCA GAA GGA GTC
CAC CTT ACA TCT TGT GCT AAG GCT CCG CG-3'

5
1R
5'-ATA CCT TCC TTG TCT TGA ATT TTC GCC TTA ACG TTC TCG
ATG GTA TCG GAA GGT TCA ACT TCC AAT GTT ATG GTT TTA
CCG GTC AAA GTC TTG ACG AAA ATC TGC ATG-3'

10
2R
5'-GAT CCG CGG AGC CTT AGC ACA AGA TGT AAG GTG GAC
TCC TTC TGA ATG TTG TAA TCA GAC AGC GTT CTA CCG TCT
TCT AGC TGC TTG CCG GCA AAG ATC AAT CTT TGT TGA TCT
GGA GGG-3'

15
3
5'-GAT CCC CGC GGT GGT GGT CCG GAA ACC CTG TGC GGT
GCT GAA CTG GTT GAC GCT CTT CAG TTC GTT TGC GGT GAC
CGT GGT TTC TAC TTC AAC AAA CCG ACC GGT TAC GGT TCC
TCC TCC CGT CGT GCT CCG CAG-3'

20
4
5'-ACC GGT ATC GTT GAC GAA TGC TGC TTC CGG TCC TGC
GAC CTG CGT CGT CTG GAA ATG TAC TGC GCT CCG CTG AAA
CCG GCT AAA TCC GCT TAA TGC ATC GAT CTC GAG-3'

25
3R
5'-AGC ACG ACG GGA GGA GGA ACC GTA ACC GGT CGG TTT
GTT GAA GTA GAA ACC ACG GTC ACC GCA AAC GAA CTG
AAG AGC GTC AAC CAG TTC AGC ACC GCA CAG GGT TTC CGG
ACC ACC ACC GCG GG-3'

30
4R
5'-AAT TCT CGA GAT CGA TGC ATT AAG CGG ATT TAG CCG
GTT TCA GCG GAG CGC AGT ACA TTT CCA GAC GAC GCA GGT
CGC AGG ACC GGA AGC AGC ATT CGT CAA CGA TAC CGG TCT
GCG G-3'

35

40 The ubiquitin and IGF-I sequences were isolated from pPO39354 and
pPO39334 (by SphI-SacII and SacII-NsiI digests, respectively), and cloned into
SphI-NsiI digested pDM25454 (a pUC19-based plasmid carrying a sequence coding for
DsbA), to create a plasmid, designated pPO39358, containing a DsbA::ubiquitin::IGF-I
fusion gene.

The fusion gene from pPO39358 was ligated into the double-cassette binary parent vectors pDM25470 and pDM25465 to create pPO39377 and pPO41623, respectively. EcoRI-XbaI fragments of pPO39377 and pPO41623 were ligated to form the chromosomal transfer DNA (Figure 17).

- 5 The chromosomal transfer DNA was transformed into *E. coli* strain B1384, which contains an attP site as well as a sequence, under the control of the trp promoter, encoding the enzyme integrase (INT). Indole acrylic acid (1 mM) was added to induce the expression of INT and resulted in the integration of transduced chromosomal transfer DNAs. Cells were tested for chromosomal transfer DNA
- 10 integration by:

Blue/yellow screening Cells were tested for integrated DNA by blue/yellow screening with AmpScreen (BRL). Colonies with a blue phenotype were further screened, yellow colonies were discarded.

PCR Cells were tested for properly integrated DNA by amplification of

15 host cell chromosomal DNA using primer pairs:

T7F1 5'-AAT TGT CGA CAT TAA TAC GAC TCA CTA TAG GGA
GAC CAC AAC GGT TTC CCT GAA TTG TCG ACA TTA ATA CGA
CTC ACT ATA GGG AGA CCA CAA CGG TTT CCC TG-3'

- 20 IGFREV 5'-CCC ATC GAT GCA TTA AGC GGA TTT AGC CGG TTT
CAG-3'

which confirm the presence of the complete fusion gene with its promoter and

- 25 T7REV 5'-TGC TAG TTA TTG CTC AGC GG-3'
- TRPBR2 5'-AAG GGC TTC ATC ATC GGT AAT AGA CA-3'

which confirm the integration of the chromosomal transfer DNA into the att site of B1384.

- 30 Production of protein from integrated genes requires T7 RNA polymerase activity, which is lacking in B1384. To test protein production from the integrated gene, P1 lysates were made using a B1384 integrant. The lysates were then transduced into *E. coli* strain W3110DE3 (as described in Example 1), which is Gal⁺ and carries a copy of the T7 RNA polymerase gene under the control of the lac promoter. Transductants were

selected by plating on galactose minimal medium plates which contained 10 1g/ml kanamycin. Single kan^r/Gal⁺ colonies were isolated and reselected on galactose minimal medium plates with kanamycin. Kan^r/Gal⁺ colonies were further analyzed by PCR using primer pairs:

5 ATT3 5'-GAG GTA CCA GCG CGG TTT GAT CAG-3'

T7RNAP1 5'-CAG CGT TAT CCG CAA CCT CAC C-3'

which showed that the upstream att site flanking the prophage in W3110DE3 is
10 unoccupied; and

T7F1 5'-AAT TGT CGA CAT TAA TAC GAC TCA CTA TAG GGA
GAC CAC AAC GGT TTC CCT GAA TTG TCG ACA TTA ATA CGA
CTC ACT ATA GGG AGA CCA CAA CGG TTT CCC TG-3'

15 IGFREV 5'-CCC ATC GAT GCA TTA AGC GGA TTT AGC CGG TTT
CAG-3'

which confirmed that the fusion gene expression cassette was transferred intact.

Individual isolates from the W3110DE3 transduction were tested for T7
20 RNA polymerase activity by streaking the isolates against phage 4107, which requires T7
RNA polymerase activity to lyse bacteria (Novagen). An isolate which contained an
intact fusion gene expression cassette and which was positive for T7 RNA polymerase
activity, designated c49222, was used to test protein production. Protein expression was
induced by the addition of IPTG to the a culture of c49222 for two hours. Protein
25 production was analyzed by SDS-PAGE of a whole cell lysate on a 12.5% acrylamide
gel. Densitometric analysis of and SDS-PAGE gel showed that the
DsbA::ubiquitin::IGF-I fusion protein accumulated to 22.3% of total cell protein.

P1 lysates were also used to transduce the integrated gene into *E. coli*
strain cDM46809 (which is cam^r, malE deleted, and contains an attB site introduced into
30 the lac region). Transductants were selected by growth on plates containing kanamycin
and chloramphenicol. Integration into the lac region was confirmed by PCR using
primer pair:

UBI1 5'-CAG ATT TTC GTC AAG ACT TTG ACC GGT AAA ACC
ATA ACA TTG GAA GTT GAA CCT TCC GAT ACC ATC GAG AAC

GTT AAG GCG AAA ATT CAA GAC AAG GAA GGT ATC CCT
CCA GAT CA-3'

1224 5'-CGC CAG GGT TTT CCC AGT CAC GAC-3'

5

A P1 lysate was then made from an isolate which was kan^r/cam^r and integrated into the *lac* region. This P1 was used to transduce W3110DE3. Transductants were selected for kanamycin and chloramphenicol resistance by growth on selective media. Kan^r/cam^r isolates were tested for T7 RNA polymerase activity by streaking against phage 4107 as described above. Two isolates positive for T7 RNA polymerase activity, designated c49258#46 and c49258#50, were tested for protein accumulation by induction with IPTG for two hours. Whole cell lysates were analyzed by SDS-PAGE using 12.5% acrylamide gels. DsbA::ubiquitin::IGF-I fusion protein accumulated to 19.6% of total cell protein in c49258#46, as measured by densitometry of an SDS-PAGE gel.

15

Southern blot analysis of chromosomal DNA from c49222 and c49258#46 was performed to check the copy number of the integrated DNA. Chromosomal DNA from c49222 and c49258#46 was isolated, digested with restriction endonucleases, transferred to Hybond-N (Amersham), and probed with the a DNA fragment encoding the ubiquitin and IGF-I portions of the fusion protein. Analysis of the Southern blot showed that there were approximately two copies each of the DsbA::ubiquitin::IGF-I gene integrated into the chromosomes c49222 and c49258#46 (Figure 24), i.e. a single copy of the integrated DNA). This result was surprising and unexpected in view of the levels of accumulation of DsbA::ubiquitin::IGF-I protein shown by SDS-PAGE (22.3% and 19.6% of total cell protein, respectively). Ordinarily, it is expected that such high levels of protein accumulation can only be accomplished by expression of heterologous genes carried by high copy number plasmids.

20

DsbA::ubiquitin::IGF-I was also produced by integrating a chromosomal transfer DNA carrying a gene for tetracycline resistance in addition to the gene for kanamycin resistance. P1 lysates prepared from a B1384 integrant were used to transduce W3110DE3 to kanamycin resistance (see Example 1). Kan^r isolates were checked for properly integrated DNA using primer pairs T7F1 x IGFREV and ATT3 x T7RNAP1 as described above. Isolates were also tested for T7 RNA polymerase activity

30

by streaking against phage 4107 as described above. Isolates positive for T7 RNA polymerase activity were then selected for amplification of the integrated DNA by growth on medium containing kanamycin (10 µg/ml) and tetracycline (30µg/ml). The tetracycline allele incorporated into this construct is effective at high copy number, therefore colonies which are tetracycline resistant may have amplified the integrated DNA. Kan^r/tet^r colonies were tested for protein accumulation by induction with IPTG, as described above. All kan^r/tet^r colonies produced the fusion protein upon induction.

Example 3

10 Chromosomal expression of ubiquitin hydrolase (UBP-1)

The construction of plasmids used in this example is described in Figures 10-16. pJT70 was the source of the ubiquitin hydrolase. pDM25493 was the source of the *trp* promoter used for this construct. chromosomal transfer DNA's for the yeast UB-1 gene under the control of the *trp* promoter were prepared from pDM46813 and either pDM25472 or pDM25448. In this example, pDM25472 was used (i.e. chromosomal transfer DNA#1 of Figure 16). The fusion gene formed by this chromosomal transfer DNA encodes an in-frame fusion between a truncated DsbA gene and a UB-1 cDNA missing the amino-terminal 92 codons.

The chromosomal transfer DNA was introduced into B1384 as in Example 2. Integrants were selected for with kanamycin (10 µg/ml). Isolated colonies were tested in a diagnostic PCR reaction using primers TRPPF and 1239 (as described in Example 1). All isolates were positive by this test. All isolates were also ampicillin sensitive.

One colony was selected for further characterization. P1 lysates were prepared of this isolate and used to transduce W3110DE3 to kanamycin resistance as described in Example 1. Kanamycin resistant colonies were further tested by PCR using primers ATT3 and T7RNAP1, as described in Example 2. All isolates showed the expected location at the *attB/P* or *attP/B* sites flanking the DE3 lysogen.

The isolates were tested for protein expression by testing for ubiquitin hydrolase activity. Isolates were grown in casamino acid minimal medium, harvested and lysed by sonication. The soluble fraction was assayed for activity by incubation with

DsbA::ubiquitin::IGF-I fusion protein substrate at 37° C for one hour. Cleavage was monitored by SDS-PAGE. All isolates (WBD311, 312, 313, 314, 331, and 332) showed good levels of enzyme activity (i.e. complete cleavage of the substrate under assay conditions).

5

Example 4

Expression of an insulin-like growth factor binding protein-3 (IGFBP-3) fusion protein

A chromosomal transfer DNA carrying a fusion protein comprising DsbA, a linker including a human rhinovirus 2A protease site, and IGFBP-3

10 (DsbA::2A::IGFBP-3) was created using the double cassette method. Construction of the fusion gene and chromosomal transfer DNA are shown in Figure 18. DsbA was from pDM46905, the 2A protease site was created by annealing primers V2ATA and V2ATB, and IGFBP-3 was PCR amplified from pYZ42580 using primers BP3RZ and NBP3F.

15 The IGFBP-3 gene used to create the DsbA::2A::IGFBP-3 fusion was created by annealing and ligating a number of synthetic oligonucleotides, which, when fully assembled, code for IGFBP-3 protein. The oligonucleotides were assembled in three segments; 5', 3', and middle. Oligonucleotides

20 F1-1 5'-AGC TTG GTG CTT CTT CTG CTG GTC TTG GAC CAG
TTG TTC GTT GTG AAC CAT GTG ATG CAC GAG CTT TAG CTC
AAT GTG CTC CAC CAC CAG CTG TT-3',

25 F1-2 5'-TGT GCT GAA TTA GTT CGA GAA CCA GGT TGT GGT
TGT TGT TTA ACT TGT GCT TTA TCT GAA GGT CAA CCA TGT
GGT ATT TAT ACT GAA CGT TGC GG-3',

30 F1-3 5'-TAG TGG TTT GCG TTG TCA ACC AAG CCC AGA TGA
AGC TAG GCC TTT ACA AGC ATT ATT AGA TGG TCG AGG TCT
GTG TGT TAA TGC GTC CGC TGT TTC TCG ATT GCG CGC G-3',

35 C1-1 5'-TCG ACG CGC GCA ATC GAG AAA CAG CGG ACG CAT
TAA CAC ACA GAC CTC GAC CAT CTA ATA ATG CTT GTA AAG
GCC TAG CTT CAT CTG GGC TTG GTT G-3',

C1-2 5'-ACA ACG CAA ACC ACT ACC GCA ACG TTC AGT ATA
AAT ACC ACA TGG TTG ACC TTC AGA TAA AGC ACA AGT TAA
ACA ACA ACC ACA ACC TGG TTC TC-3',

and

5 C1-3 5'-GAA CTA ATT CAG CAC AAA CAG CTG GTG GTG GAG
CAC ATT GAG CTA AAG CTC GTG CAT CAC ATG GTT CAC AAC
GAA CAA CTG GTC CAA GAC CAG CAG AAG AAG CAC C-3'

were annealed and ligated to form the 5' segment of the IGFBP-3, then cloned into
pUC18 (HindIII-Sall digest); this construct was designated pYZ37437. The 3' section
of the gene was created by annealing and ligating oligonucleotides

10 F-1 5'-TCG ACG TGA GAT GGA GGA TAC CTT AAA CCA TTT
AAA ATT TTT GAA CGT TTT ATC CCC GCG TGG CGT TCA TAT
CCC GAA TTG CGA T-3',

15 F-2 5'AAA AAA GGC TTC TAC AAA AAG AAA CAA TGC CGT
CCG AGT AAG GGT CGT AAA CGA GGT TTT TGT TGG TGC GTT
GAC AAA TAC GGT-3',

20 F-3 5'-CAA CCG TTG CCG GGT TAT ACT ACT AAA GGC AAA
GAA GAT GTT CAT TGT TAT TCT ATG CAA TCT AAA TAA TGC
ATC TCG AG-3',

25 C-1 5'-AAT TCT CGA GAT GCA TTA TTT AGA TTG CAT AGA
ATA ACA ATG AAC ATC TTC TTT GCC TTT AGT AGT ATA ACC
CGG C-3',

C-2 5'-AAC GGT TGA CCG TAT TTG TCA ACG CAC CAA CAA
AAA CCT CGT TTA CGA CCC TTA CTC GGA CGG CAT TGT TTC
TTT TTG TAG AAG-3',

30 and

35 C-3 5'-CCT TTT TTA TCG CAA TTC GGG ATA TGA ACG CCA
CGC GGG GAT AAA ACG TTC AAA AAT TTT AAA TGG TTT AAG
GTA TCC TCC ATC TCA CG-3',

followed by cloning into Sall-EcoRI digested pUC18 (designated pYZ37405).

pYZ374100 contained the middle segment of the IGFBP-3 gene and was created by
annealing and ligating oligonucleotides

40 MF1 5'-CGC GCT TAT TTA TTA CCT GCC CCA CCG GCA CCG
GGT AAC GCC TCC GAA A-3',

MF2 5'-GCG AAG AGG ATC GTT CTG CGG GTT CCG TTG AAT
CTC CAA GTG TGA GTT CTA CCC ATC GAG TTA GCG ACC CGA
AA-3',

5 MF3 5'-TTT CAT CCG TTG CAC TCT AAA ATC ATT ATT ATT
AAA AAG GGT CAC GCA AAG GAT TCT CAA CGT TAT AAG
GT-3',

10 MF4 5'-GGA TTA TGA AAG CCA ATC TAC CGA CAC TCA AAA
TTT TAG TAG TGA AAG TAA ACG TGA AAC CGA GTA CGG CCC
GTG-3',

15 MB1 5'-TCG ACA CGG GCC GTA CTC GGT TTC ACG TTT ACT
TTC ACT ACT AA-3',

MB2 5'-AAT TTT GAG TGT CGG TAG ATT GGC TTT CAT AAT
CCA CCT TAT AAC GTT GAG AAT CCT TTG CGT GAC CCT TTT
T-3',

20 MB3 5'-AAT AAT AAT GAT TTT AGA GTG CAA CGG ATG AAA
TTT CGG GTC GCT AAC TCG ATG GGT AGA ACT CAC ACT TGG
AGA TT-3',

and

25 MB4 5'-CAA CGG AAC CCG CAG AAC GAT CCT CTT CGC TTT
CGG AGG CGT TAC CCG GTG CCG GTG GGG CAG GTA ATA
AAT AAG-3',

digesting the ligated DNA with BssHII and SalI, end filling with Klenow then cloning
into Klenow-filled, XbaI-digested pUC18.

30 PCR amplification of a segment of pYZ37490 was used to add a SacII site
and repair a cloning artifact. Primer pairs

pF1 5'-GGT TGT TGT TTA ACT TGT GCT TTA TCT GAA GGT
CAA CCA TGT GGT ATT TAT ACT GAA CGT TGC GGT AGT GGT
TTG CGT TGT CAA CCA AGC CCA GAT GAA GCT AGG-3'

35

1233 5'-AGC GGA TAA CAA TTT CAC ACA GGA-3'

and

pR1 5'-TAA AGC ACA AGT TAA ACA ACA ACC ACA ACC TGG
TTC TCG AAC TAA TTC AGC ACA AAC AGC TGG TGG TGG AGC
ACA TTG AGC TAA AGC TCG TGC ATC ACA TGG T-3'
1224 5'-CGC CAG GGT TTT CCC AGT CAC GAC-3'

40

were used to introduce the restriction site and repair the defect. The two PCR amplified fragments were then mixed and amplified to form a single DNA using primer pair 1233 x 1224. The resulting DNA segment was cloned into HindIII-Sall digested pYZ37437, creating pYZ37490.

5 PCR amplification was also used to introduce an additional SacII site, to facilitate later cloning steps. Primers

5pMP 5'-GAC TGC AAG CTT CCG CGG TGG TGG TGC TTC TTC
TGC TGG TCT TGG A-3'

and

10 1233 5'-AGC GGA TAA CAA TTT CAC ACA GGA-3'

were used to amplify a segment of pYZ37490, which was then ligated into HindIII-Sall digested pYZ37490, forming pYZ42519.

The IGFBP-3 gene was assembled from the three segments in a three-way
15 ligation reaction. pYZ42519 (HindIII-BssHII digest), pYZ374100 (BssHII-Sall digest) and pYZ37405 (Sall-EcoRI digest) were ligated into HindIII-EcoRI digested pUC18. A properly assembled clone was identified by restriction mapping and sequencing.

Cloning artifacts were repaired using PCR. BPFIX1 was created by
amplifying pYZ42509 with primers

20 YZM1 5'-CTC GAT TGC GCG CTT ATT TAT TAC C-3'

and

YZM2 5'-TCT CAC GTC GAC ACG GGC CGT ACT CGG TTT CAC
GTT TAC TCA GTA CTA AAA T-3',

25 and cloning the resulting fragment (BssHII-Sall digested) into a BssHII-Sall digest of pYZ42509. A HindIII-BssHII digest of BPFIX1 was ligated with a HindIII-BssHII digest of pYZ42519 to create pYZ42529. A second repair was made using primer pairs

715F1' 5'-TGT TGG TGC GTC GAC AAA TAC GGT C-3'

30 1233 5'-AGC GGA TAA CAA TTT CAC ACA GGA-3'

and

715R' 5'-GAC CGT ATT TGT CGA CGC ACC AAC A-3'

1224 5'-CGC CAG GGT TTT CCC AGT CAC GAC-3'.

35

This repaired a cloning defect and added a Sall site. Two DNA fragments were amplified from pYZ42529 using primer pairs 715F1' x 1233 and 715R' x 1224. These two fragments were mixed and PCR amplified into a single DNA fragment using 1233 x 1224. This single fragment was digested with BssHI and Sall, then ligated into a

5 BssHI-Sall digest of pYZ42529, creating pYZ50559.

pYZ42580, the donor construct for the IGFBP-3 gene, was created by ligation of EcoRI-SacII fragments from pYZ50559 and pDM25497.

The chromosomal transfer DNA carrying the DsbA::2A::IGFBP-3 fusion gene were transfected into *E. coli* strain B1384, which was grown in the presence of 100

10 1M IAA to induce the expression of INT and the integration of the chromosomal transfer DNA. Integrants were selected with kanamycin. All isolates were also ampicillin sensitive.

Isolates were further characterized by diagnostic PCR amplification of the host cell chromosome. PCR amplification with primer pairs

15 1227 5'-TAA TAC GAC TCA CTA TAG GGA GA-3'
BP3-607 5'-GGG ATA TGA ACG CCA CGC GGG GAT AA-3',

INT107 5'-GCG GAG AAA CCA TAA TTG CAT CTA CTC-3'
BP3-559 5'-CGT GAA ACC GAG TAC GGC CCG TGT C-3',

20 and

T7REV 5'-TGC TAG TTA TTG CTC AGC GG-3'
TRPBR2 5'-AAG GGC TTC ATC ATC GGT AAT AGA CA-3'

confirmed the proper integration of the intact chromosomal transfer DNA into the

25 chromosome at the att site.

P1 lysates were prepared from a single isolate and used to transduce W3110DE3 to kanamycin resistance (as described in Example 1). Kanamycin resistant isolates were assayed for T7 RNA polymerase activity by streaking against phage 4107, as described in Example 2. Isolates with T7 RNA polymerase activity were then tested

30 for expression of the fusion gene by induction with IPTG, followed by analysis of protein expression by SDS-PAGE of whole cell lysates on 12.5% polyacrylamide gels. Densitometric analysis of whole cell lysates indicated that the DsbA::2A::IGFBP-3 fusion protein accumulated to a level of 22.6% (Figure 25 A).

Example 5

Production IGF-I fusion proteins

Fusion proteins including DsbA and IGF-I linked by a sequence including a site for either human rhinovirus 2A or 3C protease were produced using the double
5 cassette binary system. Construction of binary plasmids and chromosomal transfer DNAs is diagramed in Figures 19 and 20.

For expression of DsbA::2A::IGF-I, EcoRI/XbaI digest fragments of pPO53096 and pPO57211 were ligated to form the chromosomal transfer DNA (CTD-DsbA::2A::IGF-I). EcoRI/XbaI fragments of pPO53097 and pPO57210 were
10 ligated to form a chromosomal transfer DNA carrying a gene encoding DsbA::3C::IGF-I (CTD-DsbA::3C::IGF-I). CTD-DsbA::3C::IGF-I and CTD-DsbA::2A::IGF-I were each transformed into B1384 cells in the presence of indole acrylic acid (to induce INT expression). Transformants were grown on media containing kanamycin to select for integrants. Nine individual kan^r colonies from each transformation were tested for
15 ampicillin sensitivity. All tested colonies were ampicillin sensitive.

Isolates were tested for correctly integrated DNA by PCR amplification with primer pairs T7F1 x IGFREV and T7REV x TRPBR2 to confirm the presence of the intact fusion gene and integration into the att site of B1384, as described in Example 2.

P1 lysates were prepared from one of the B1384 integrants from each
20 transformation and used to transduce W3110DE3 to kanamycin resistance. Kan^r/gal⁺ isolates were tested for the presence of T7 RNA polymerase activity as described in Example 2. Isolates positive for T7 RNA polymerase activity were further tested by PCR using primer pairs T7F1 x IGFREV and ATT3 x T7RNAP1 to confirm appropriate integration of the intact fusion gene, as described in Example 2.

25 Two isolates from each transduction (c57265#44 and c57265#54 for DsbA::2A::IGF-I; c57264#5 and c57264#28 for DsbA::3C::IGF-I) were then grown on medium containing both kanamycin and tetracycline. Both DsbA::3C::IGF-I and CTD-DsbA::2A::IGF-I carry a tetracycline resistance allele which confers resistance when the gene is in high copy number. Growth in the presence of tetracycline selects for
30 amplification of the integrated DNA. Both isolates from each transduction were kan^r/tet^r. The isolates were then tested for expression of the fusion proteins by induction with

IPTG. Protein expression was assayed by SDS-PAGE of whole cell lysates.

Densitometric scanning of a SDS-PAGE gel showed that the two isolates expressing DsbA::3C::IGF-I fusion protein accumulated the fusion protein to 20% and 20.1% of total cell protein and the two isolates expressing DsbA::2A::IGF-I accumulated the fusion protein to 25.7% and 38% of total cell protein.

Example 6

Chromosomal expression of TGF- β 2 using the double cassette binary system

A chromosomal transfer DNA encoding a fusion protein comprising DsbA, ubiquitin, and human TGF-J2 (DsbA::ubiquitin::TGF- β 2) was created using the double cassette method. Construction of the fusion gene and chromosomal transfer DNA are shown in Figure 21. DsbA::ubiquitin was from pDM25497, and TGF-J2 was PCR amplified from pPC-21 (Madisen et. al. (1988) DNA 7:1-8) using primers

UBTGFJ2F 5'-GGG GCC GCG GTG GTG CTT TGG ATG CGG CCT
ATT GCT TTA GA-3'
and

TGFJ2R 5'-GGG GAA TTC TTA GCT GCA TTT GCA AGA CTT TAC
A-3'.

pDM25497 was digested with SacII-EcoRI and the 4.3 kb fragment containing pUC18 and DsbA::ubiquitin sequences was isolated. The 0.35 kb PCR product resulting from the amplification of pPC-21 encoding the last 112 amino acids of human TGF-J2 was purified and digested with SacII-EcoRI. These two fragments were ligated to create pDP26, a pUC18 derivative containing a DsbA::ubiquitin::TGF-J2 fusion gene. pDP26 was the donor construct for assembly of the binary plasmids used to make the chromosomal transfer DNA.

The fusion gene from pDP26 was ligated into the double-cassette binary vectors pDM25470 and pDM25465 to create pC9DP and pA6DP, respectively. Briefly, pDM 25470 was digested with BamHI-SmaI and the 4.2 kb fragment was isolated. pDP26 was digested with EcoRI, blunt ended with the Klenow fragment of DNA polymerase, and then digested with BamHI. The 1.1 kb fragment from this digest was isolated. The two fragments described above were ligated to create pC9DP.

pDM25465 was digested with BamHI, blunt ended with Klenow, digested with XbaI and the 7.1 kb fragment was isolated. pDP26 was digested with EcoRI, blunt ended with Klenow, digested with BamHI, and the 1.2 kb fragment was isolated. This 1.2 kb fragment was ligated to the 7.1 kb fragment from pDM25465 to create pA6DP.

5 An additional binary plasmid containing a tetracycline resistance selectable marker was created using the 7.2 kb fragment isolated from pDM46932 following XbaI-XhoI digestion, and the fusion gene from pA6DP (2.7 kb XbaI-XhoI fragment). These two fragments were ligated to create pA6DPIIT. EcoRI-XbaI fragments of pC9DP (2.2 kb) and pA6DPIIT (6.4 kb) were ligated to form the chromosomal transfer DNA (Figure 21).

10 The chromosomal transfer DNA was transformed into *E. coli* strain B1384, which was grown in the presence of 500 μ M IAA to induce the expression of INT and the integration of the chromosomal transfer DNA. Integrants were selected with 10 μ g/ml kanamycin. All isolates were found to be ampicillin sensitive.

15 Isolates were further characterized by diagnostic PCR amplification of host cell chromosomal DNA. PCR amplifications with primer pairs

1227 5'-TAA TAC GAC TCA CTA TAG GGA GA-3'
β21079 5'-GGA AAT GGA TAC ACG AAC CC-3',
and

20 INT107 5'-GCG GAG AAA CCA TAA TTG CAT CTA CTC-3'
6HEβ2 5' GGG GGA TCC GAT CGT GGA GGA TGA TTA AAT GCA
CCA CCA CCA CCA CCA CGA CGA CGA CAA AGC TTT GGA
TGC GGC CTA T-3'

25 and primers T7REV and TRPBR2, described previously (see Example 2), confirmed the proper integration of the intact chromosomal transfer DNA into the chromosome at the att site.

 P1 lysates were prepared from a single isolate and used to transduce W3110DE3 to kanamycin resistance, as described previously. Amplification of the
30 integrated fusion gene is accomplished by growth of kanamycin resistant isolates on medium containing kanamycin and tetracycline (30 μ g/ml). Kanamycin/tetracycline resistant isolates were assayed for T7 RNA polymerase activity by streaking against phage 4107, as described in Example 2. Isolates with T7 RNA polymerase activity were then tested for expression of the fusion gene by induction with IPTG, followed by

analysis of protein expression by SDS-PAGE of whole cell lysates on 10% polyacrylamide gels (Figure 26). Protein accumulation in chromosomal integrants was comparable to the levels seen in host cells containing a multicopy number plasmid utilizing the same T7 promoter linked to the a copy of the gene encoding the

5 DsbA::ubiquitin::TGF- β 2 fusion protein. Densitometric analysis showed that protein accumulation in chromosomal integrants was as high as 36.7% of total cell protein.

Example 7

Expression of a heterologous protein using a promoter-less CTD

10 This example shows the use of a chromosomal transfer DNA which does not carry a promoter. The chromosomal transfer DNA carries a segment of DNA homologous to a bacterial gene (in this example, lacZ or DsbA) linked in-frame to a DNA sequence encoding a heterologous protein of interest (in this case the DsbA::3C::IGF-I fusion protein of Example 5), as well as selectable marker genes. The

15 homologous DNA encodes the 5' region of the bacterial gene. The chromosomal transfer DNA is introduced into the host cell, where it integrates into the homologous gene on the chromosome of the host cell, forming an operable linkage between the homologous gene's promoter and the DNA sequence encoding the heterologous protein of interest. Integrants are selected for using the selectable markers carried on the chromosomal

20 transfer DNA. The heterologous protein of interest is expressed through the homologous gene's promoter (Figure 8).

The DNA encoding the DsbA::3C::IGF-I fusion protein is constructed as described in Example 5. This fusion gene is then placed in frame to a DNA segment encoding the first 100 amino-terminal amino acids of the lacZ gene, forming a

25 lacZ/DsbA::3C::IGF-I gene. The cyclophilin, kanamycin resistance, and tetracycline resistance genes utilized in Example 5 are also cloned into the plasmid carrying the lacZ/DsbA::3C::IGF-I gene. This plasmid is then cleaved with restriction endonucleases to remove the plasmid origin of replication, the ampicillin resistance gene and other non-essential sequences, then re-ligated to form a circular chromosomal transfer DNA.

30 The chromosomal transfer DNA is transformed into E. coli host cells and the transformed host cells are grown on media containing kanamycin (10 μ g/ml).

Kanamycin resistant isolates are tested for ampicillin sensitivity, to show that the host cells carry integrated DNA, not plasmid DNA. Kan^r isolates are also tested using PCR. PCR primers from the lacZ promoter and the DNA sequence encoding DsbA are used to confirm integration of the intact chromosomal transfer DNA. Amplification of the integrated fusion gene is selected for by growth of kan^r isolates on medium containing kanamycin and tetracycline (30 µg/ml). Expression of the integrated fusion gene is induced by growth of kan^r/tet^r isolates in the presence of IPTG. Protein expression is assayed by SDS-PAGE.

Promoter-less chromosomal transfer DNAs may also be integrated into other sites on the host cell chromosome (Figure 9). Specialized host cells may be constructed which carry a chromosomal copy of an inducible promoter (in this case the T7 promoter) linked to a particular gene (in this case DsbA). This host cell is made by transforming a variant chromosomal transfer DNA (carrying a copy of the lacZ gene, the T7 promoter operably linked to the 5' end of the DsbA gene and the chloramphenicol resistance gene) into the host cell (in this case W3110DE3, which also carries a copy of the gene encoding T7 RNA polymerase). Integration of the chromosomal transfer DNA is selected for by growth of transformed W3110DE3 cells on medium containing chloramphenicol. The integration of the chromosomal transfer DNA produces a W3110DE3 host cell containing the T7 promoter linked to the 5' portion of the DsbA gene. This integrated DNA then becomes the target for integration of a chromosomal transfer DNA carrying a DNA sequence encoding the heterologous protein of interest.

A chromosomal transfer DNA carrying the DNA sequence encoding the heterologous protein of interest is constructed (in this case the DsbA::3C::IGF-I fusion gene described above and in Example 5). The cyclophilin, kanamycin resistance and tetracycline resistance genes are also cloned onto the plasmid. This plasmid is then cleaved with the appropriate restriction enzymes to remove the plasmid origin of replication, ampicillin resistance gene, and other non-essential sequences, and re-ligated to form a circular chromosomal transfer DNA. The chromosomal transfer DNA is transformed into the T7-DsbA W3110DE3 host cells described above. Integrants are selected by growth on medium containing chloramphenicol and kanamycin. Kan^r/cam^r isolates are checked for integration of the intact chromosomal transfer DNA by PCR.

PCR amplification of host cell chromosomal DNA using primer pairs T7F1 x IGFREV confirms the integration of the intact chromosomal transfer DNA. Integrants are checked for T7 RNA polymerase activity by streaking against phage 4107, as described in Example 2. Amplification of the integrated DNA is selected for by growth of T7 RNA polymerase-positive isolates on kanamycin, chloramphenicol, and tetracycline. Resistant isolates are assayed for protein expression by induction with IPTG. Protein expression is assayed by SDS-PAGE.

Example 8

10 Expression of a DsbA::3C::IGFBP-3 fusion protein using the double cassette system

A gene encoding DsbA::3C::IGFBP-3 fusion protein was expressed using the double cassette binary system shown in Figure 7. The DsbA sequence was originally isolated by PCR amplification of the DsbA gene from the *E. coli* chromosome; plasmid pDM25454 was used as the source of the DsbA sequence for this fusion gene. The site for 3C protease was created by synthesizing two oligonucleotides, 15 RV3CTA 5'-CCCGATTCTCTGGAAGTTCTGTTCCAA-3' and RV3CTB 5'-TTGGAACAGAACTTCCAGAGAATCGGGCATG-3', which were annealed to form a double stranded DNA fragment encoding a 3C protease cleavage site. The IGFBP-3 gene was constructed by annealing and ligating synthetic oligonucleotides, as described in Example 4. The IGFBP-3 sequence used for construction of the gene encoding the DsbA::3C::IGFBP-3 fusion protein was a PCR amplified DNA fragment made using primers BP3RZ and NBP3F and template pYZ42580. Cloning of the two DNA sources used to make chromosomal transfer DNA 20 carrying the gene encoding the DsbA::3C::IGFBP-3 fusion protein, pDM46947 and pDM46948, is shown in Figure 22.

The chromosomal transfer DNA was constructed using EcoRI/XbaI fragments from pDM46947 and pDM46948. The chromosomal transfer DNA was transformed into B1384 cells grown in the presence of indole acrylic acid (to induce the expression of INT). Integrants were selected for by growth of transformants on media 30 containing kanamycin. All kanamycin resistant isolates were also ampicillin sensitive.

Kanamycin resistant isolates were checked by PCR using primer pairs 1227 x BP3-607, INT107 x BP3-559, and T7REV x TRPBR2, as described in Example 4.

- P1 lysates were prepared from one of the kanamycin resistant isolates and used to transduce W3110DE3 to kanamycin resistance. Kanamycin transductants were
- 5 tested for the presence of T7 RNA polymerase activity by streaking against phage 4107, as described in Example 2. Kanamycin resistant/T7 RNA polymerase positive isolates were selected for chromosomal amplification by growth on media containing kanamycin and tetracycline. One kan^r/tet^r isolate was selected and checked for protein expression by induction with IPTG. Protein accumulation was assayed by SDS-PAGE (Figure 25 B).
- 10 Densitometric analysis of an SDS-PAGE gel showed that the DsbA::3C::IGFBP-3 fusion protein accumulated to an average of 27.4% of total cell protein.

- All publications, patents and patent applications cited in this specification are incorporated herein by reference to the same extent as if each individual publication, patent, or patent application was specifically and individually indicated to be
- 15 incorporated by reference.

It should be apparent that one having ordinary skill in the art would be able to surmise equivalents to the claimed invention which would be within the spirit of the description above. Those equivalents are to be included within the scope of the present invention.

What is claimed is:

1. A method for producing a heterologous protein of interest, comprising the steps of:
 - 5 transferring a chromosomal transfer DNA into a bacterial host cell, wherein said chromosomal transfer DNA comprising at least one copy of a gene encoding the heterologous protein of interest and a selectable marker, and wherein said host cell comprising a chromosome; selecting for integration of said chromosomal transfer DNA into said host
 - 10 cell chromosome resulting in a host cell chromosome comprising a gene encoding a heterologous protein of interest operably linked to a promoter functional in the host cell and a selectable marker flanked by duplicate DNA; and expressing said gene, wherein said gene is at no time operably linked to a promoter functional in
 - 15 a host cell on a multicopy number plasmid vector during construction of the transfer DNA and wherein said non-bacterial protein of interest accumulates within said host cell to a level in excess of 0.1% of total cell protein.
- 20 2. The method of claim 1 wherein said chromosomal transfer DNA further comprises a promoter functional in said host cell, said promoter being operably linked to said gene encoding the heterologous protein of interest, and
3. The method of claim 1 wherein said host cell chromosome further
- 25 comprises a host cell promoter and said chromosomal transfer DNA further comprises a DNA sequence homologous to a segment of the bacterial chromosome downstream from said host cell promoter, said DNA sequence linked in-frame to said gene encoding the heterologous protein of interest, wherein integration of said chromosomal transfer DNA results in the
- 30 formation of an operable linkage between said DNA sequence and the host cell promoter.

4. The method of claim 1 wherein said non-bacterial protein of interest accumulates within said host cell to a level in excess of 1% of total cell protein.

5. The method of claim 1 wherein said heterologous protein of interest is
5 a eukaryotic protein.

6. The method of claim 1 wherein said heterologous protein of interest is a mammalian protein.

10 7. The method of claim 1, wherein each said duplicate DNA comprises said gene encoding a heterologous protein of interest linked to said promoter.

8. The method of claim 1 further comprising selecting for chromosomal amplification of said chromosomal transfer DNA following integration of said
15 chromosomal transfer DNA into the chromosome of said host cell.

9. A method for producing a chromosomal transfer DNA, comprising:
ligating a restriction fragment from each of a first plasmid vector and a second plasmid vector thereby producing said chromosomal transfer DNA,
20 said first vector comprising a gene encoding a heterologous protein of interest lacking an operably linked promoter,
said second vector comprising a promoter functional in a host cell,
wherein said chromosomal transfer DNA comprises a selectable marker,
said gene encoding a heterologous protein of interest operably linked to said promoter
25 and duplicate DNA flanking said gene and lacks an origin of replication operable in said host cell.

10. A method for producing a chromosomal transfer DNA, comprising:
ligating a restriction fragment from each of a first plasmid vector and a
30 second plasmid vector thereby producing a chromosomal transfer DNA,

said first plasmid comprising a first gene encoding a heterologous protein of interest and a first promoter functional in a host cell, and wherein said first gene and first promoter are not operably linked,

5 said second vector comprising a second gene encoding a heterologous protein of interest lacking an operably linked promoter, and a second promoter functional in a host cell,

 wherein said chromosomal transfer DNA comprises a selectable marker and lacks an origin of replication operable in said host cell and wherein said first gene is operably linked to said second promoter on the chromosomal transfer DNA and said
10 second gene is operably linked to said first promoter on said chromosomal transfer DNA.

11. A chromosomal transfer DNA, comprising:

 a gene encoding a heterologous protein of interest operably linked to a promoter functional in a host cell; and

15 a selectable marker, said selectable marker flanked by duplicate DNA, wherein said gene encoding a heterologous protein of interest is at no time operably linked to a promoter functional in a host cell on a multicopy number plasmid vector.

12. A chromosomal transfer DNA, comprising:

20 two copies of a gene encoding a heterologous protein of interest, each of said copies being operably linked to a promoter functional in a host cell; and

 a selectable marker, said selectable marker flanked by said copies of said gene encoding a heterologous protein of interest,

 wherein each of said copies of said gene are at no time operably linked to
25 a promoter functional in a host cell on a multicopy number plasmid vector.

AMENDED CLAIMS

[received by the International Bureau on 10 September 1996 (10.09.96);
original claims 1-12 replaced by new claims 1-12 (3 pages)]

1. A method for producing a heterologous protein of interest, comprising the steps of:
transferring a chromosomal transfer DNA into a bacterial host cell, wherein said chromosomal transfer DNA comprises least one copy of a gene encoding the heterologous protein of interest and a selectable marker, and wherein said host cell comprises a chromosome;
selecting for integration of said chromosomal transfer DNA into said cell chromosome resulting in a host cell chromosome comprising a gene encoding a heterologous protein of interest operably linked to a promoter functional in the host cell and a selectable marker flanked by duplicate DNA; and
expressing said gene, wherein said gene is at not time operably linked to a promoter functional in a host cell on a multicopy number plasmid vector during construction of the transfer DNA and wherein said heterologous protein of interest accumulates within said host cell to a level in excess of 0.1% of total cell protein.
2. The method of claim 1 wherein said chromosomal transfer DNA further comprises a promoter functional in said host cell, said promoter being operably linked to said gene encoding the heterologous protein of interest, and wherein the operable linkage is created by circularization of the chromosomal transfer DNA.
3. The method of claim 1 wherein said host cell chromosome further comprises a host cell promoter and said chromosomal transfer DNA further comprises a DNA sequence homologous to a segment of the host cell chromosome downstream from said host cell promoter, said DNA sequence linked in-frame to said gene encoding the heterologous protein of interest, wherein integration of said chromosomal transfer DNA results in the formation of an operable linkage between said DNA sequence and the host cell promoter.
4. The method of claim 1 wherein said heterologous protein of interest accumulates within said host cell to a level in excess of 1% of total cell protein.

5. The method of claim 1 wherein said heterologous protein of interest is a eukaryotic protein.
6. The method of claim 1 wherein said heterologous protein of interest is a mammalian protein.
7. The method of claim 1 wherein each said duplicate DNA comprises said gene encoding a heterologous protein of interest linked to said promoter.
8. The method of claim 1 further comprising selecting for chromosomal amplification of said chromosomal transfer DNA following integration of said chromosomal transfer DNA into the chromosome of said host cell.
9. A method for producing a chromosomal transfer DNA comprising:
ligating a restriction fragment from each of a first plasmid vector and a second plasmid vector thereby producing said chromosomal transfer DNA, said first vector comprising a gene encoding a heterologous protein of interest lacking an operably linked promoter, said second vector comprising a promoter functional in a host cell, wherein said chromosomal transfer DNA comprises a selectable marker, said gene encoding a heterologous protein of interest operably linked to said promoter and duplicate DNA flanking said gene and lacks an origin of replication operable in said host cell.

10. A method for producing chromosomal transfer DNA comprising:

ligating a restriction fragment from each of a first plasmid vector and a second plasmid vector thereby producing a chromosomal transfer DNA, said first plasmid comprising a first gene encoding a heterologous protein of interest and a first promoter functional in a host cell, and wherein said first gene and first promoter are not operably linked, said second vector comprising a second gene encoding a heterologous protein of interest lacking an operably linked promoter, and a second promoter functional in a host cell, wherein said chromosomal transfer DNA comprises a selectable marker and lacks an origin of replication in said host cell and wherein said first gene is operably linked to said second promoter on the chromosomal transfer DNA and said second gene is operably linked to said first promoter on said chromosomal transfer DNA.

11. A chromosomal transfer DNA comprising:

a gene encoding a heterologous protein of interest operably linked to a promoter functional in a host cell; and

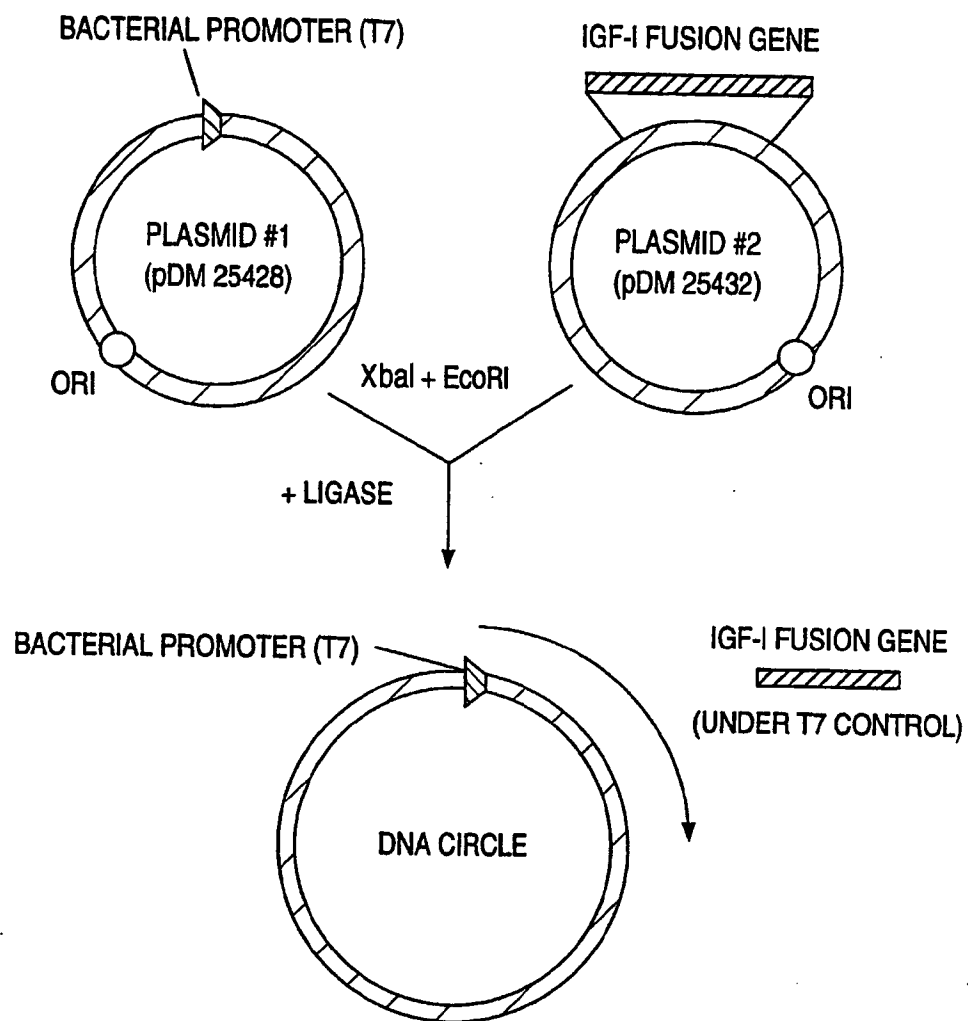
a selectable marker, said selectable marker flanked by duplicate DNA, wherein said gene encoding a heterologous protein of interest is at no time operably linked to a promoter functional in a host cell on a multicopy number plasmid vector.

12. A chromosomal transfer DNA comprising:

two copies of a gene encoding a heterologous protein of interest, each of said copies being operably linked to a promoter functional in a host cell; and

a selectable marker, said selectable marker flanked by said copies of said gene encoding a heterologous protein of interest, wherein each of said copies of said gene are at no time operably linked to a promoter functional in a host cell on a multicopy number plasmid vector.

1/26

**FIG. 1**

2/26

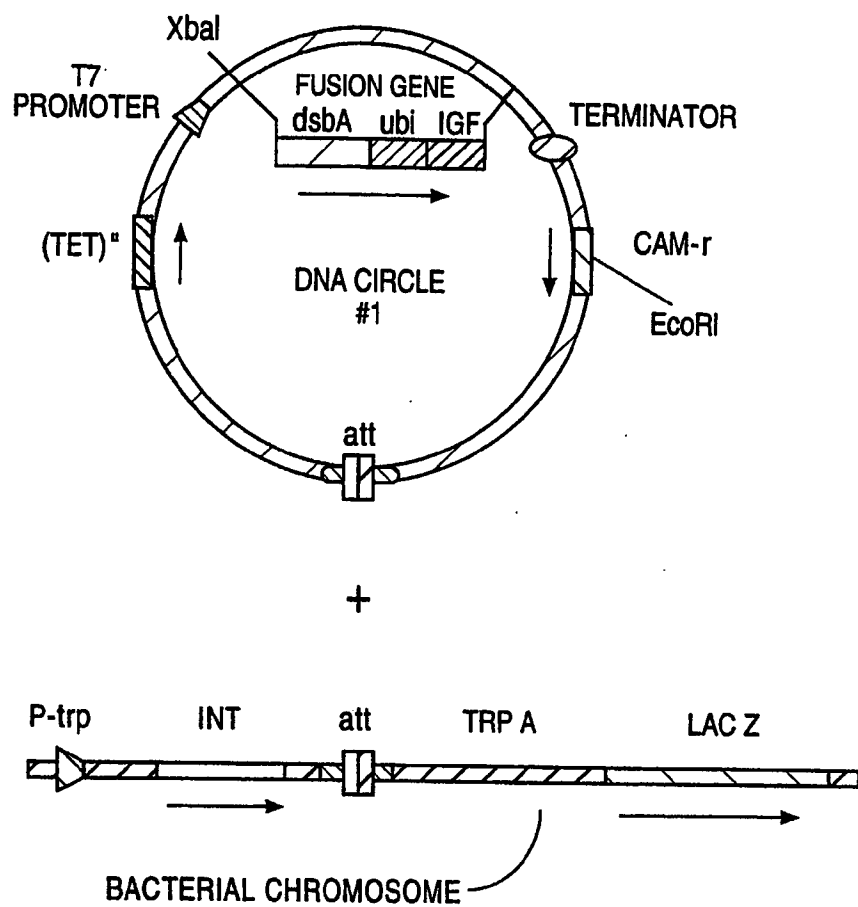


FIG. 2

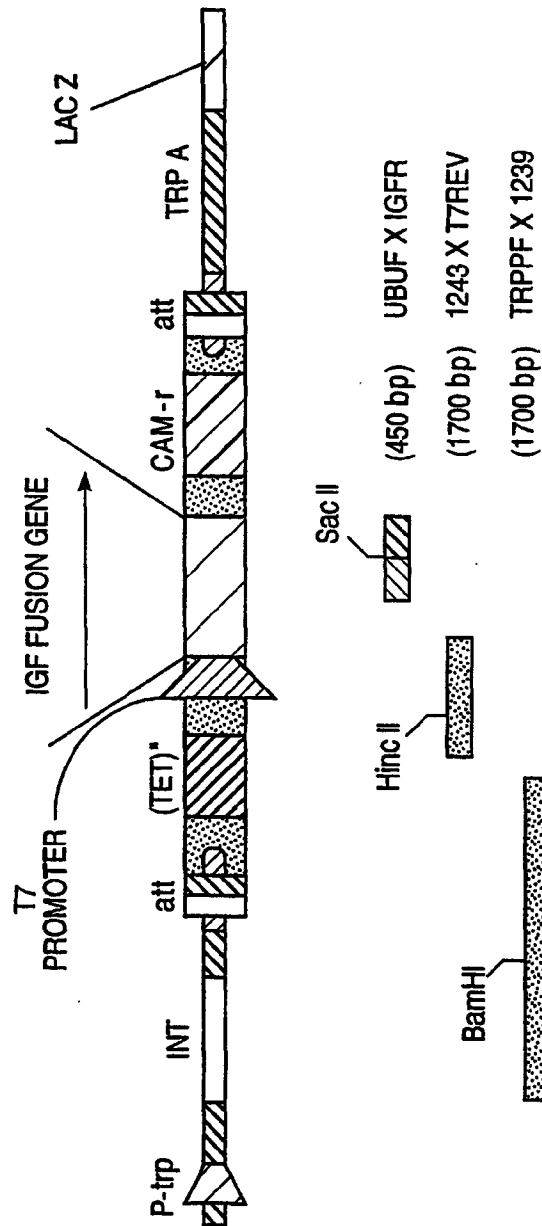
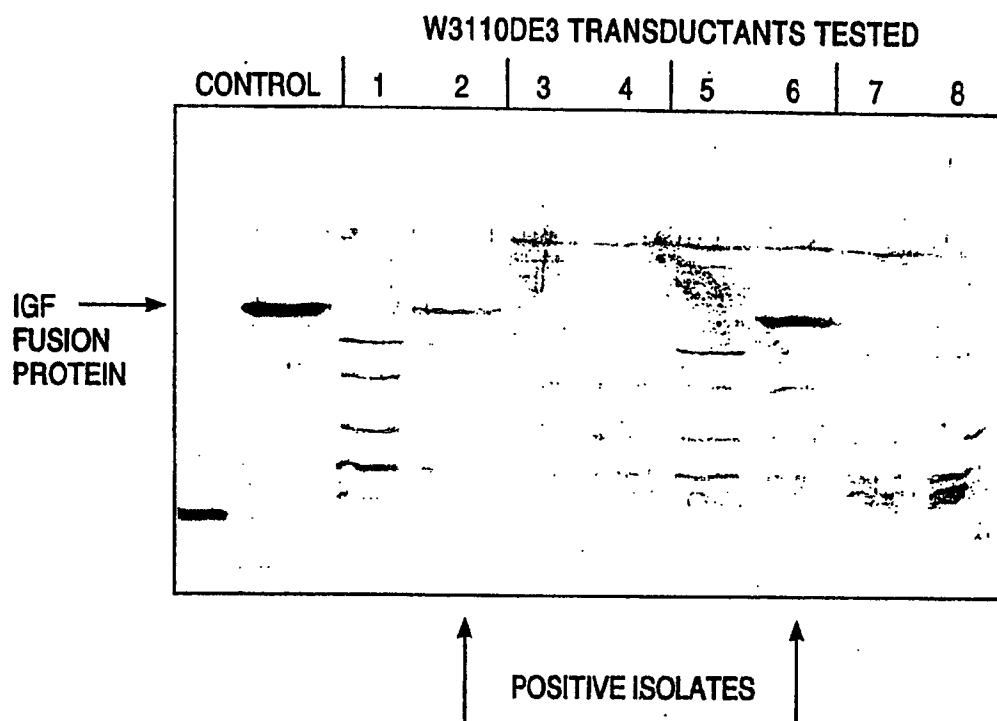


FIG. 3

4/26

**FIG. 4**

5/26

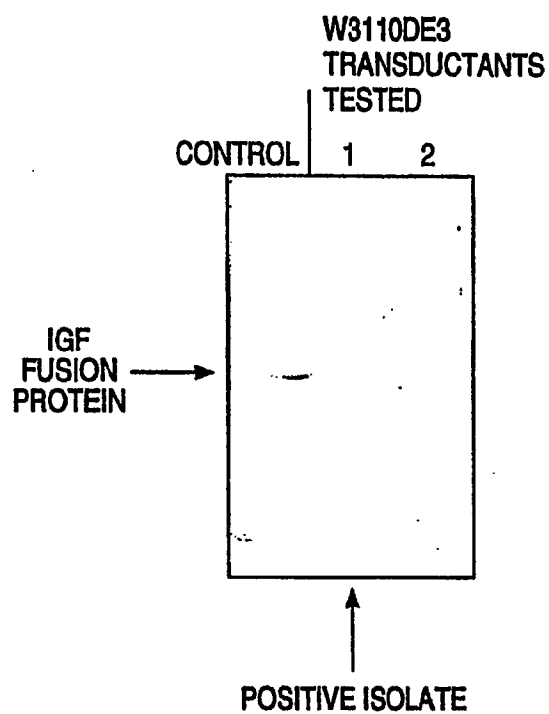
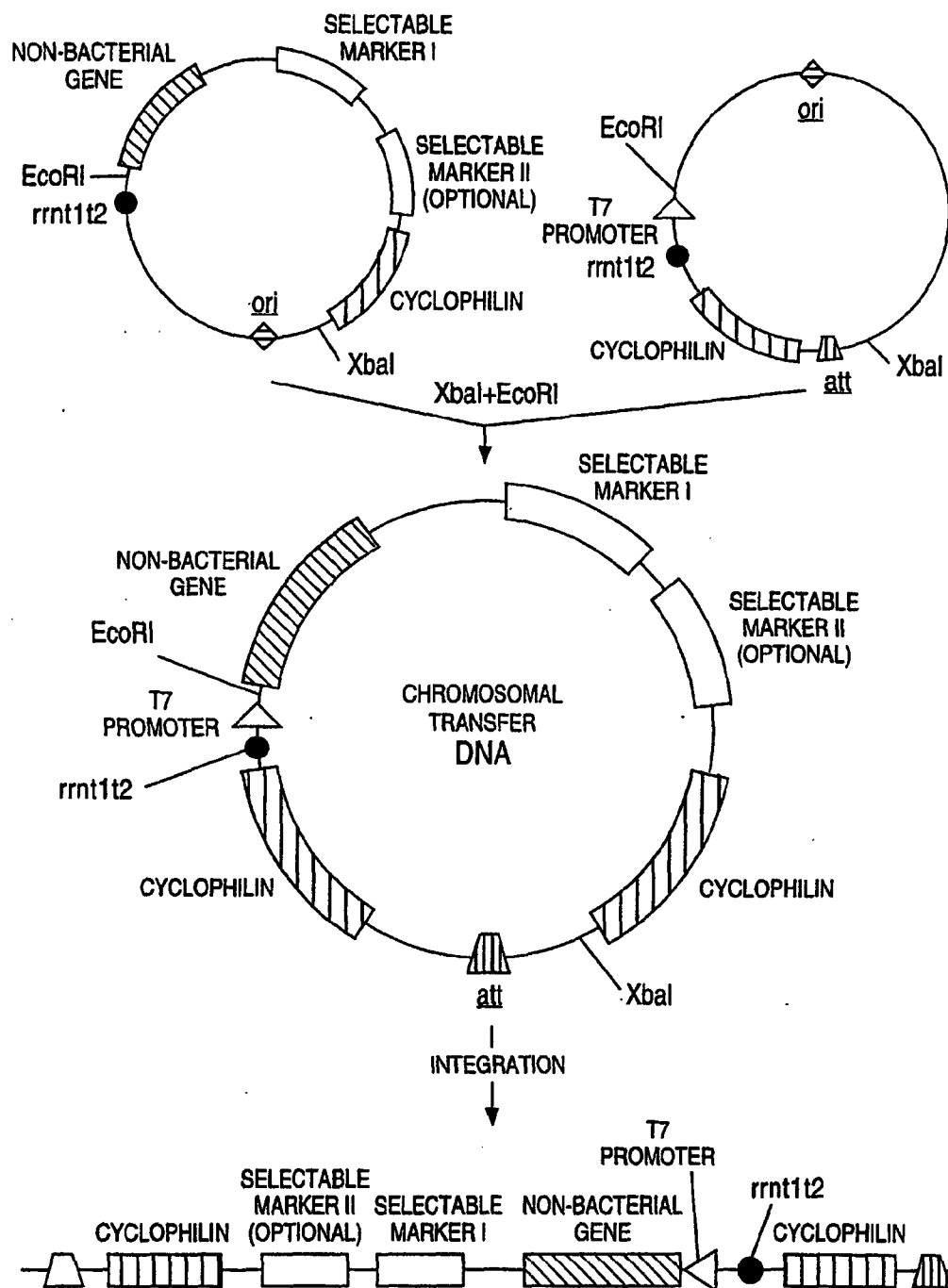


FIG. 5

6/26

**FIG. 6**

SUBSTITUTE SHEET (RULE 26)

7/26

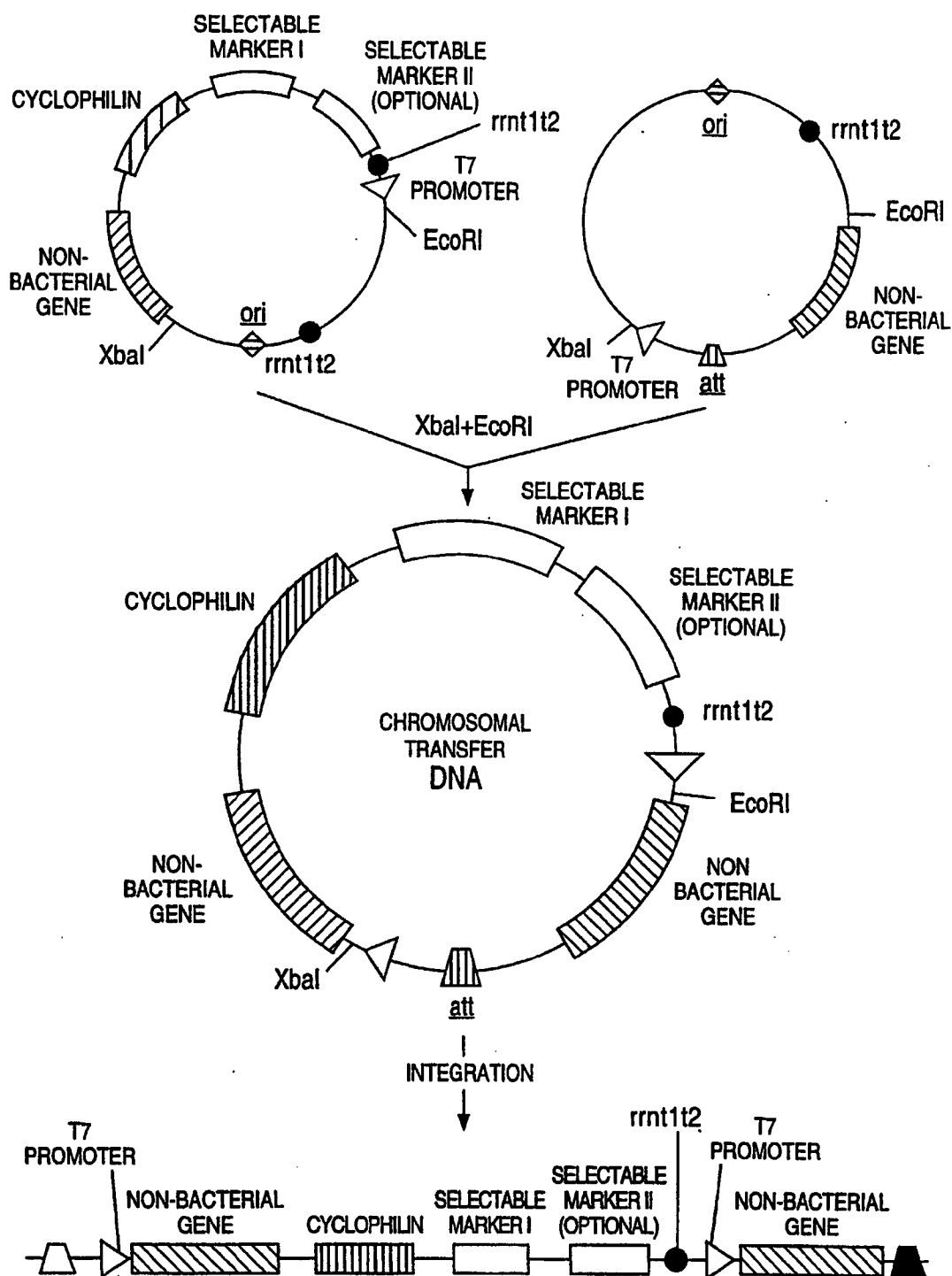
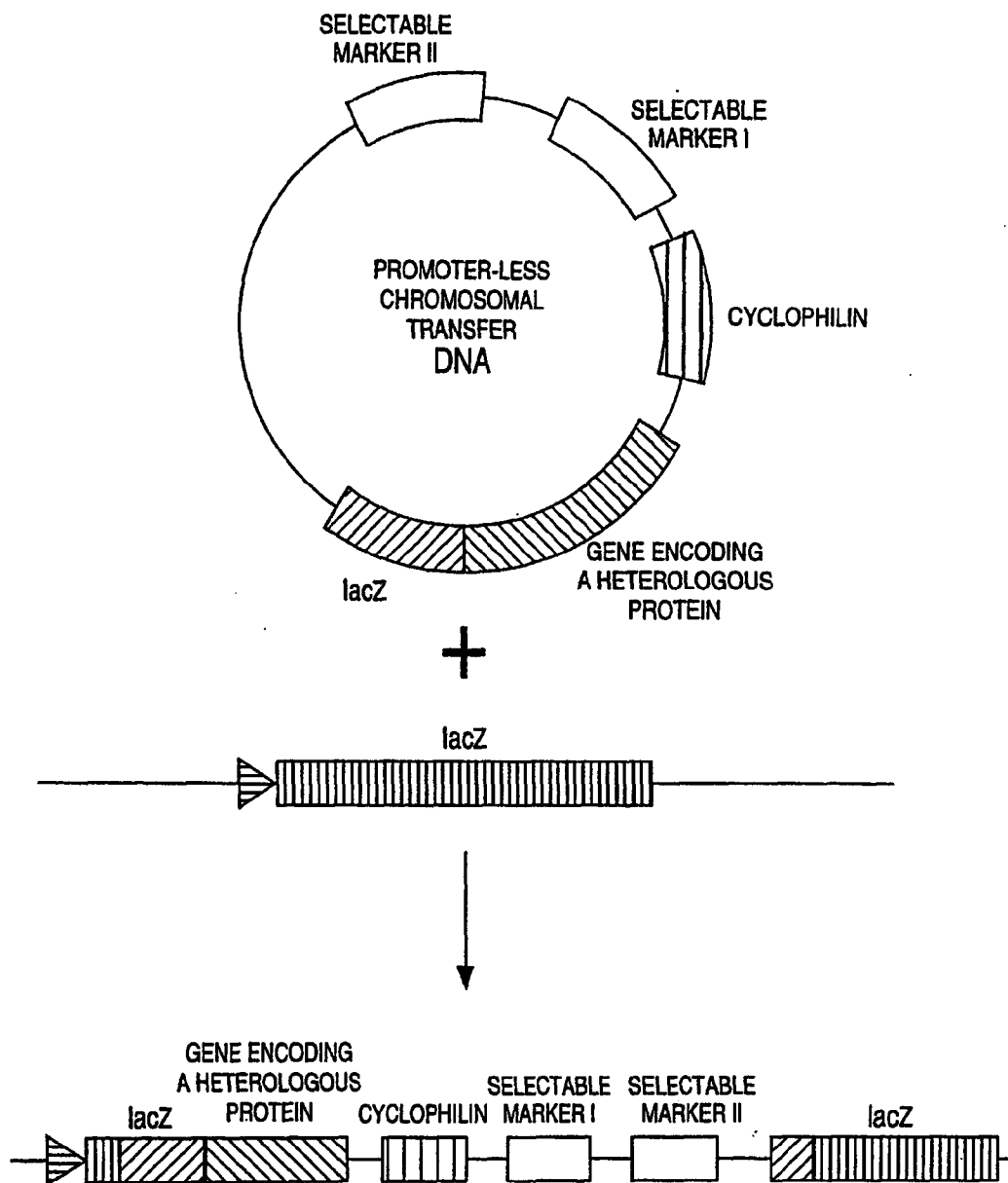


FIG. 7
SUBSTITUTE SHEET (RULE 26)

8/26

**FIG. 8**

9/26

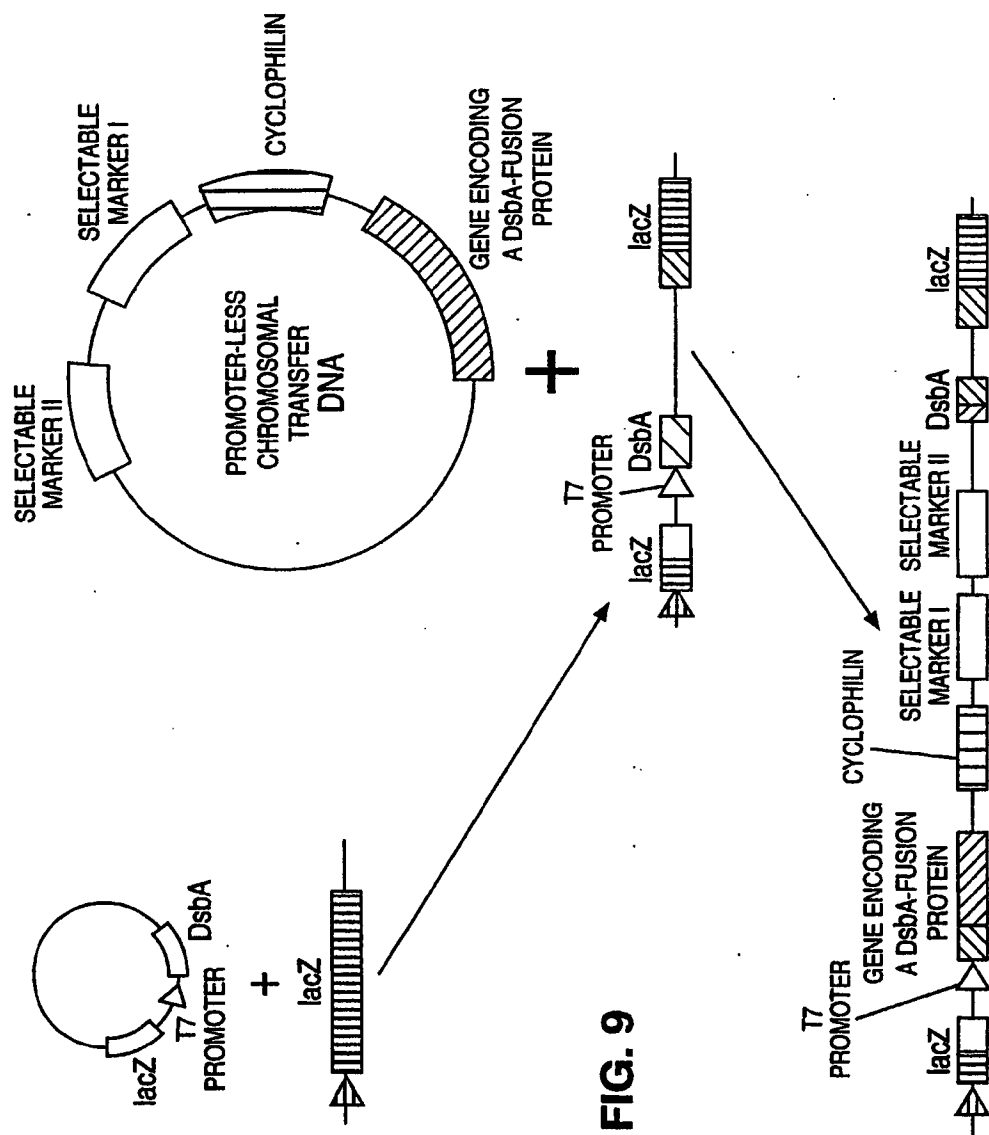


FIG. 9

10/26

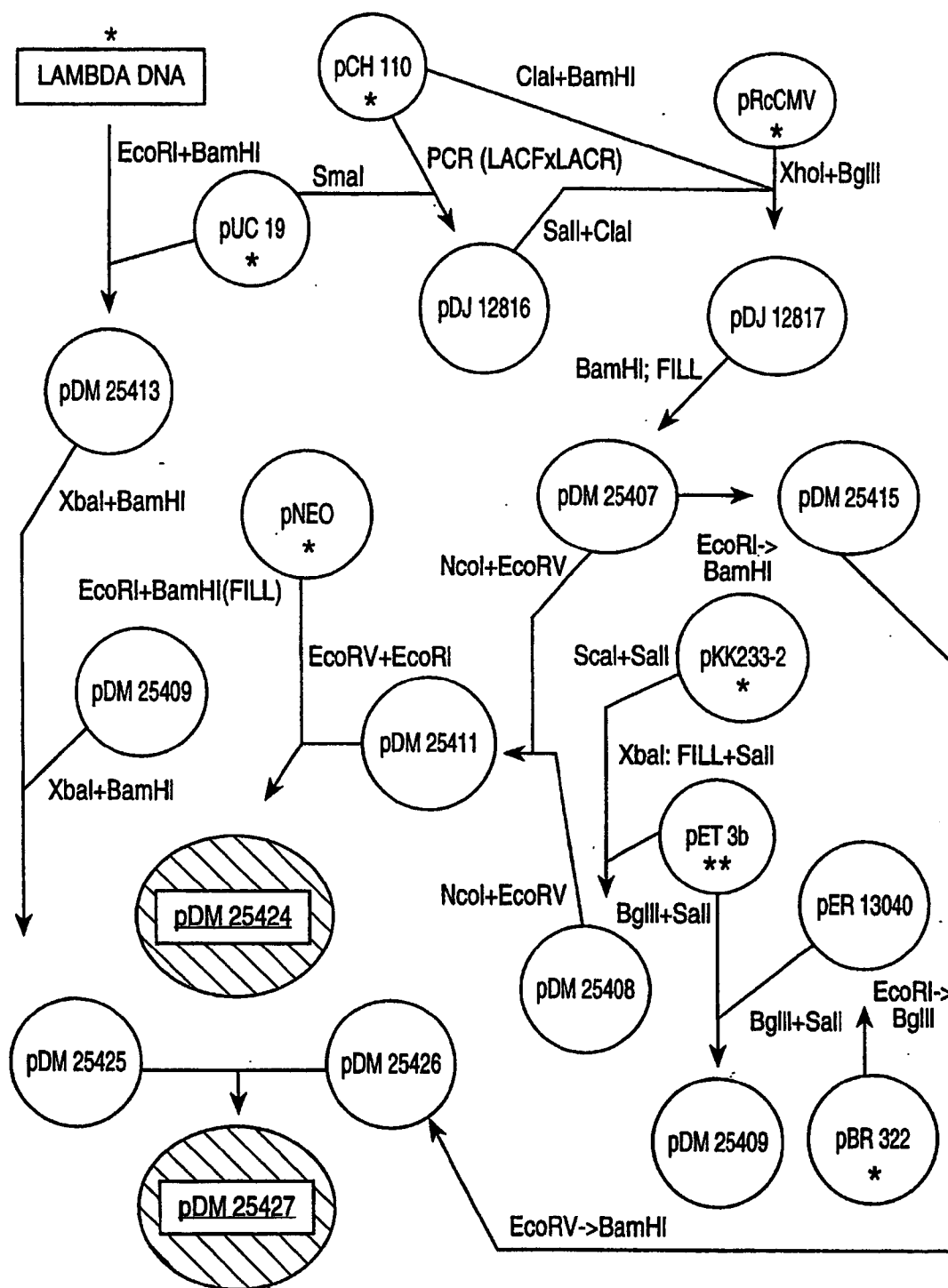


FIG. 10
RECTIFIED SHEET (RULE 91)

11/26

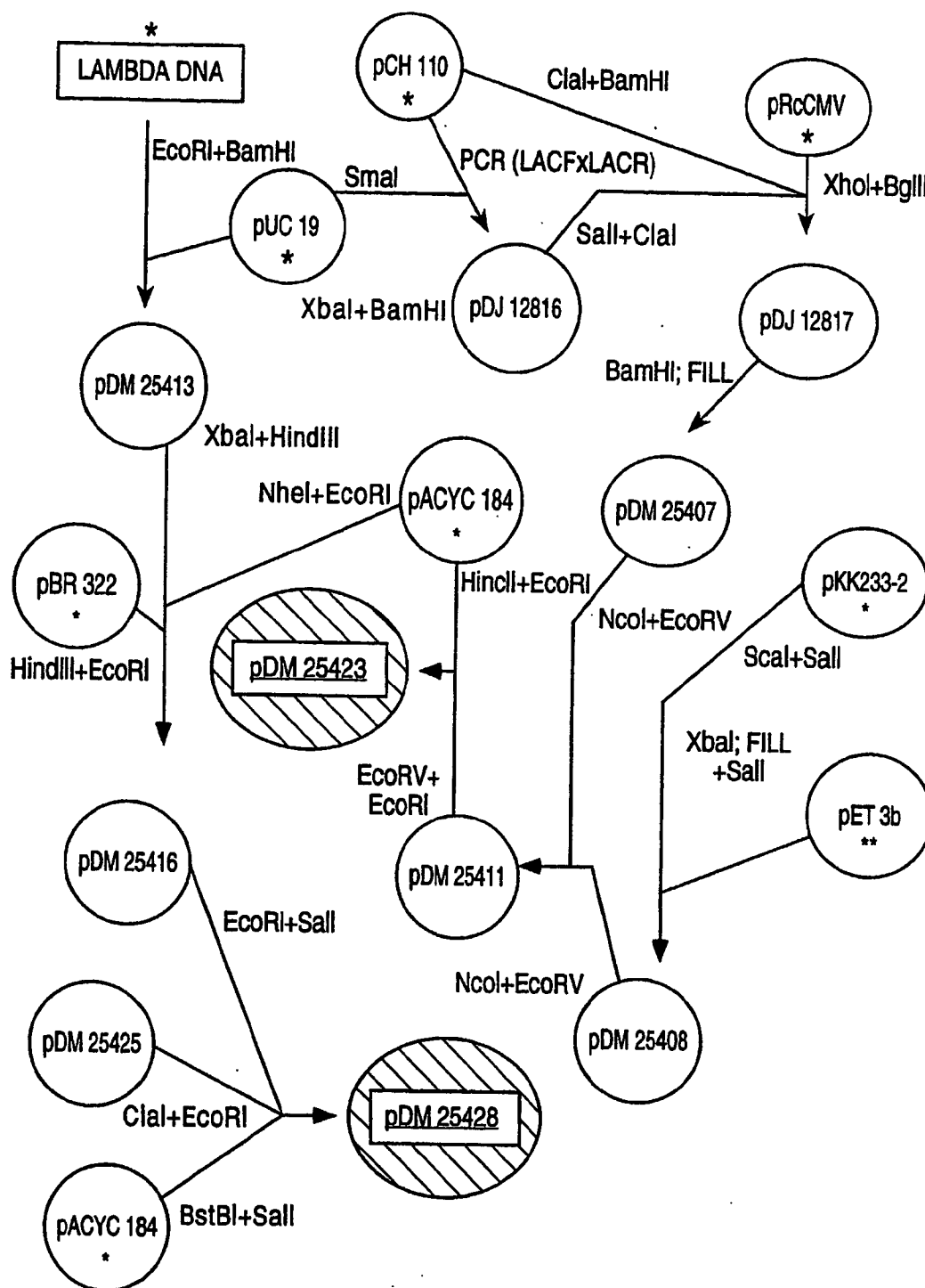


FIG. 11
RECTIFIED SHEET (RULE 91)

12/26

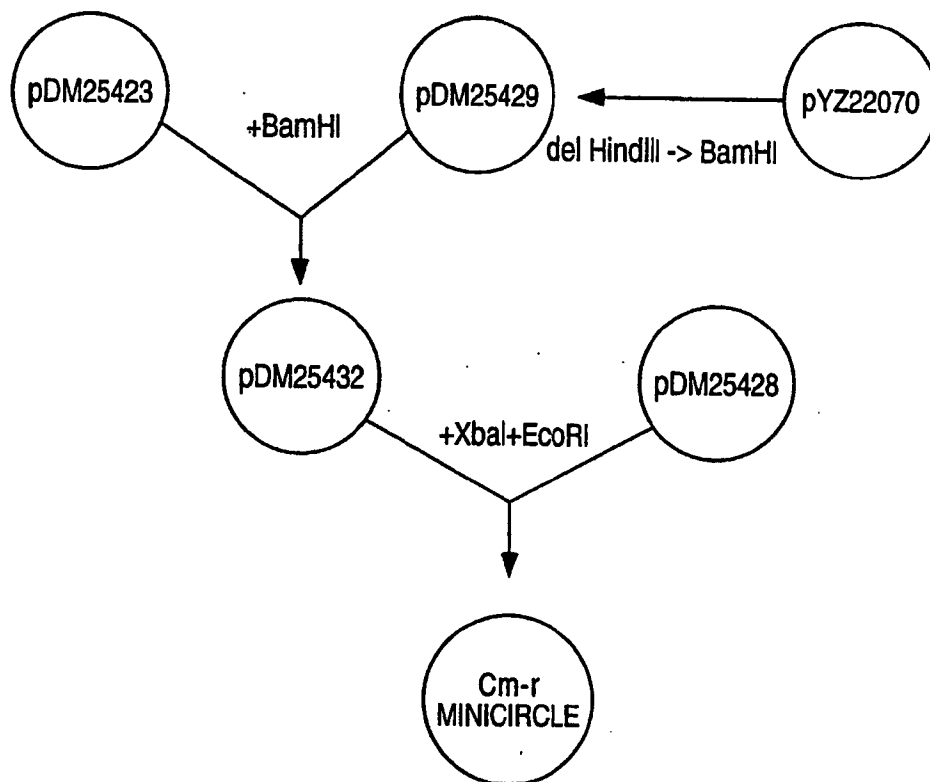
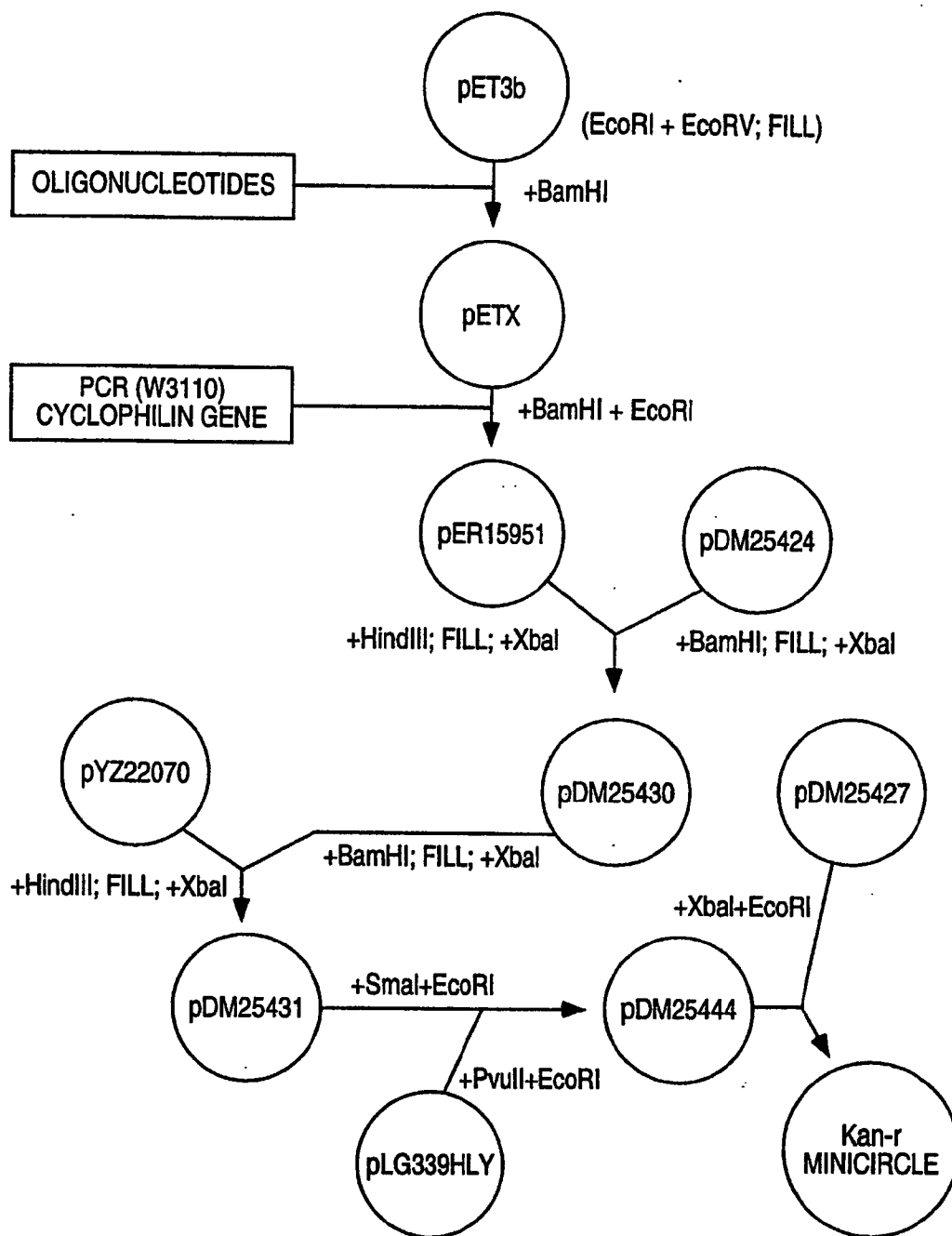


FIG. 12

13/26

**FIG. 13**

RECTIFIED SHEET (RULE 91)

14/26

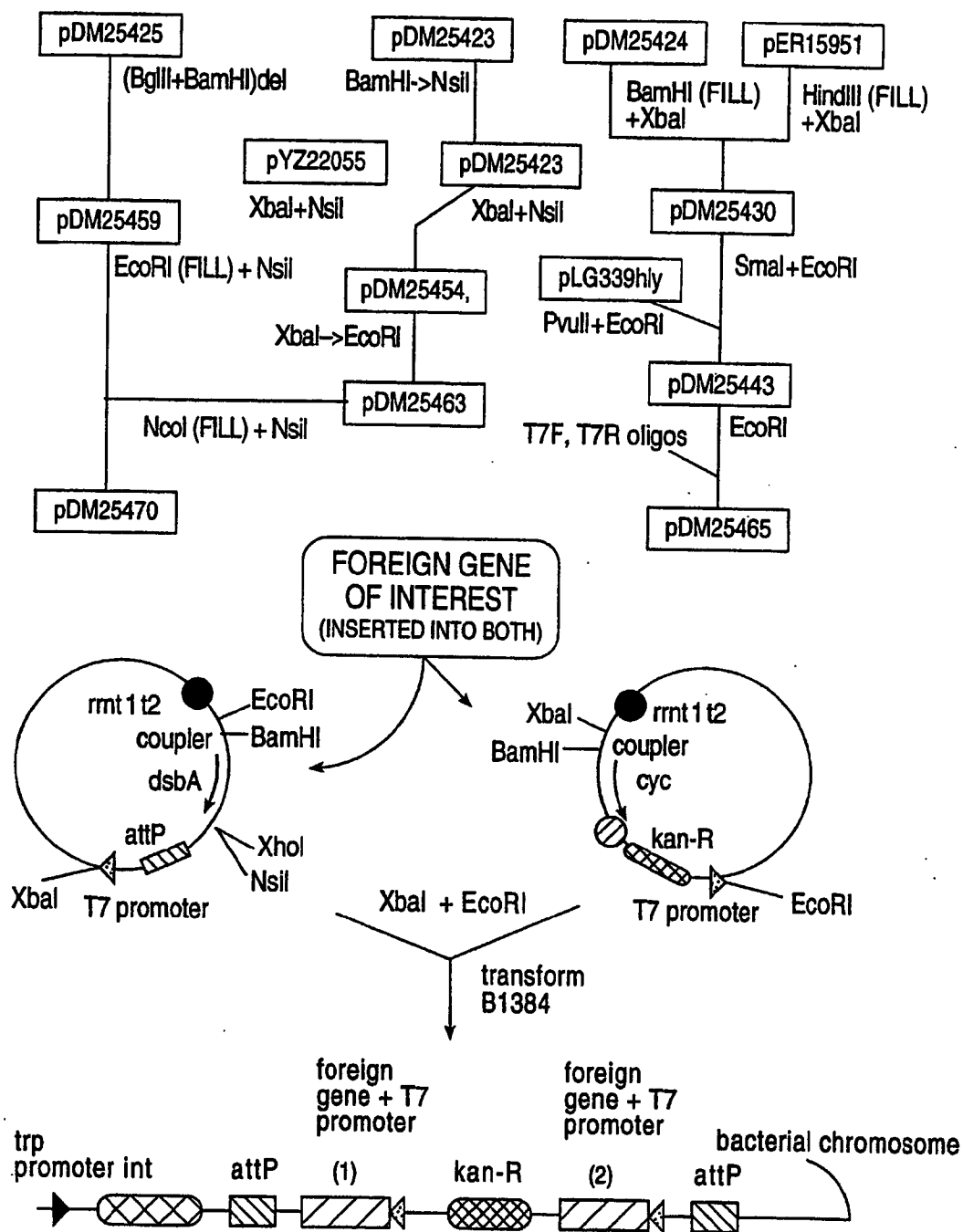


FIG. 14

RECTIFIED SHEET (RULE 91)

15/26

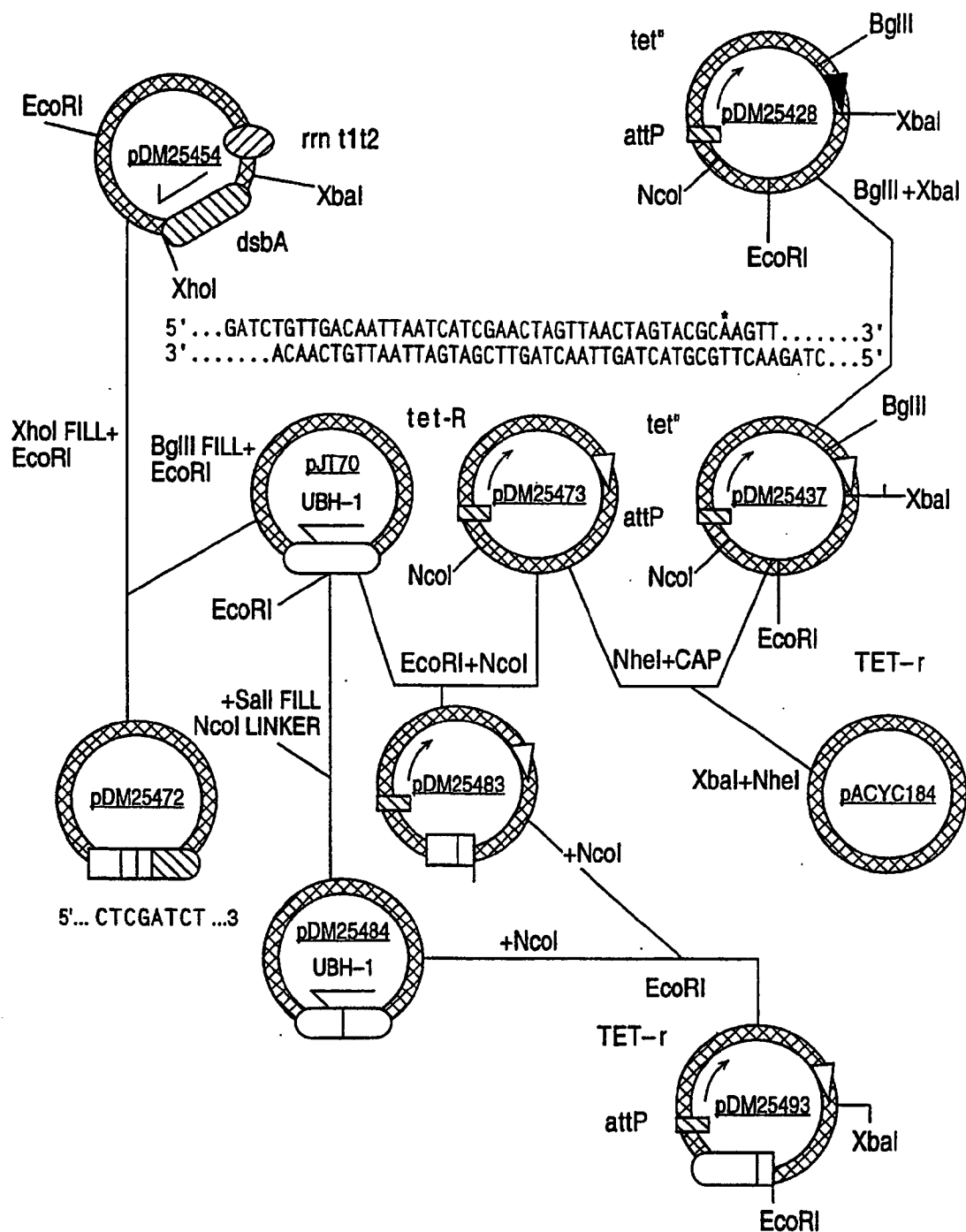


FIG. 15

SUBSTITUTE SHEET (RULE 26)

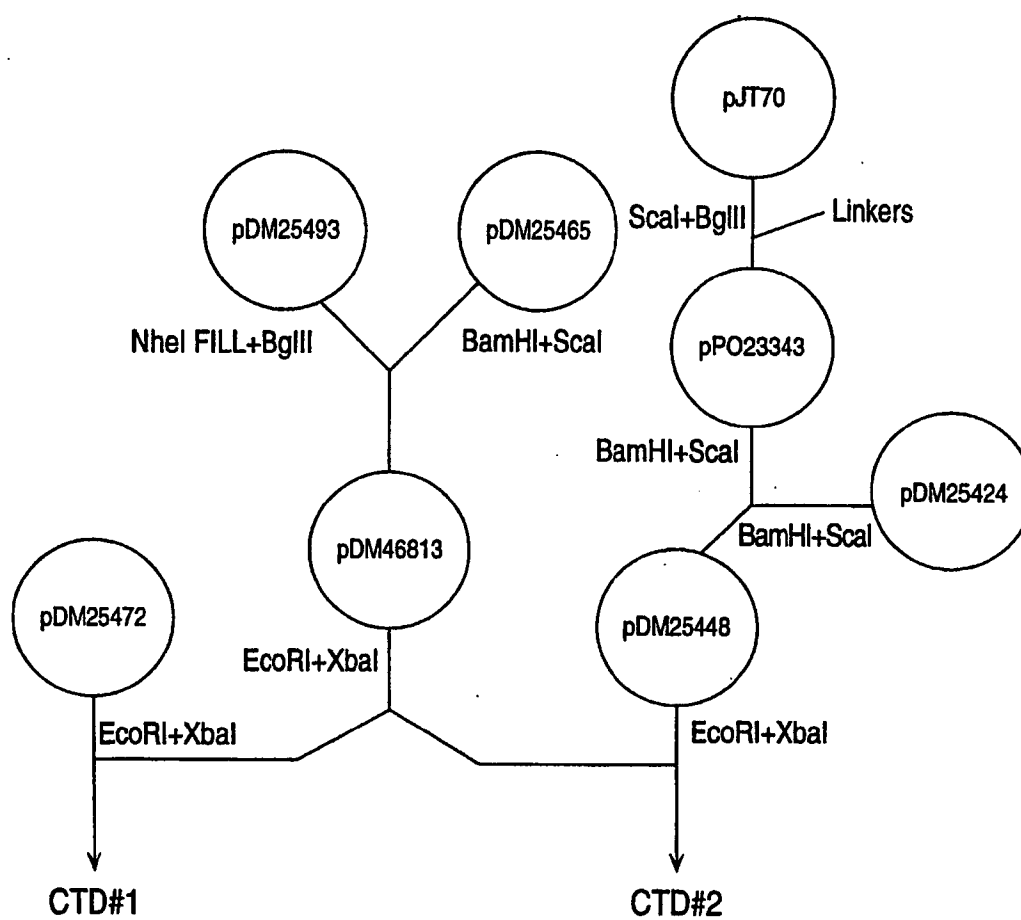


FIG. 16

17/26

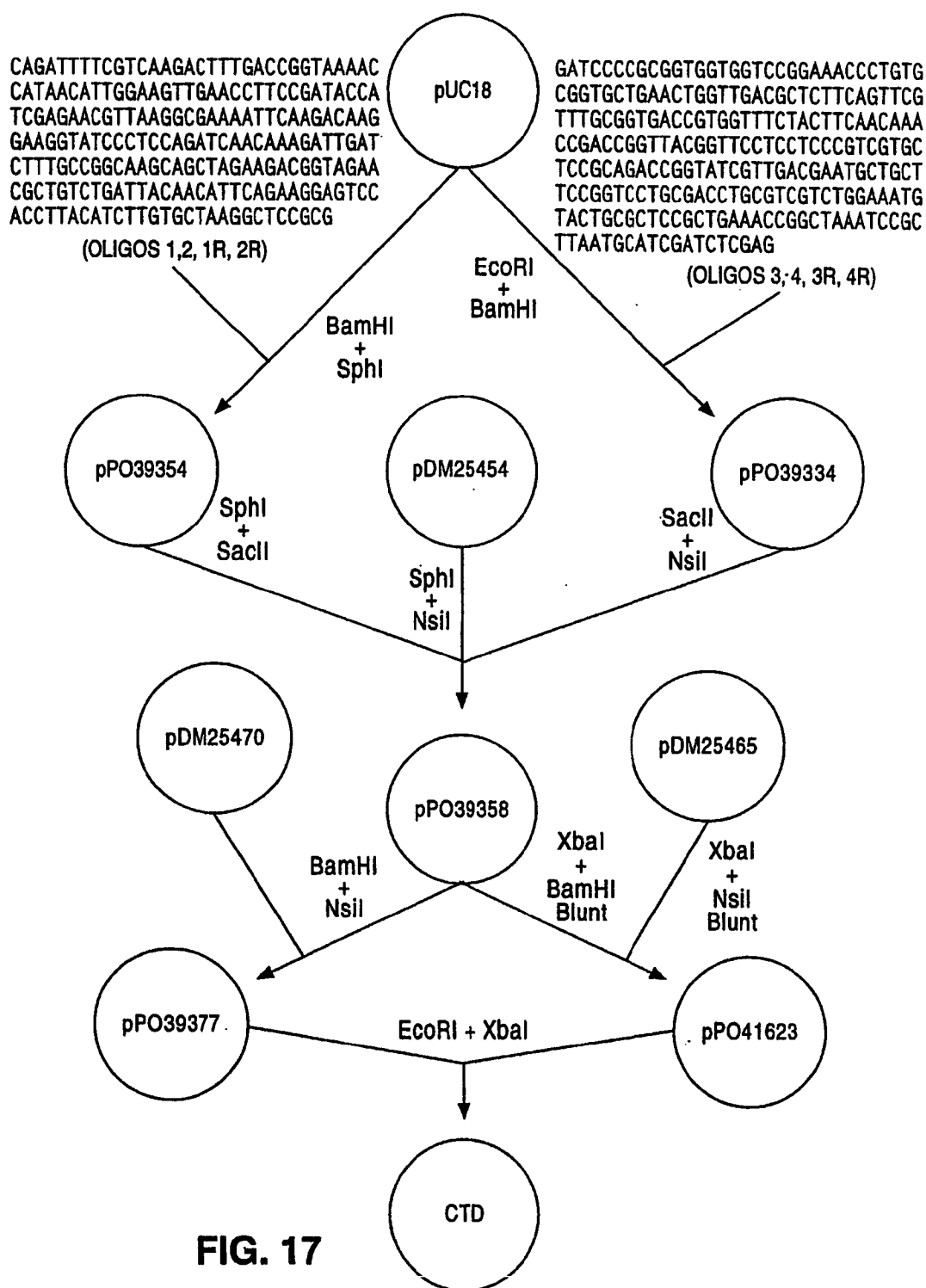


FIG. 17

18/26

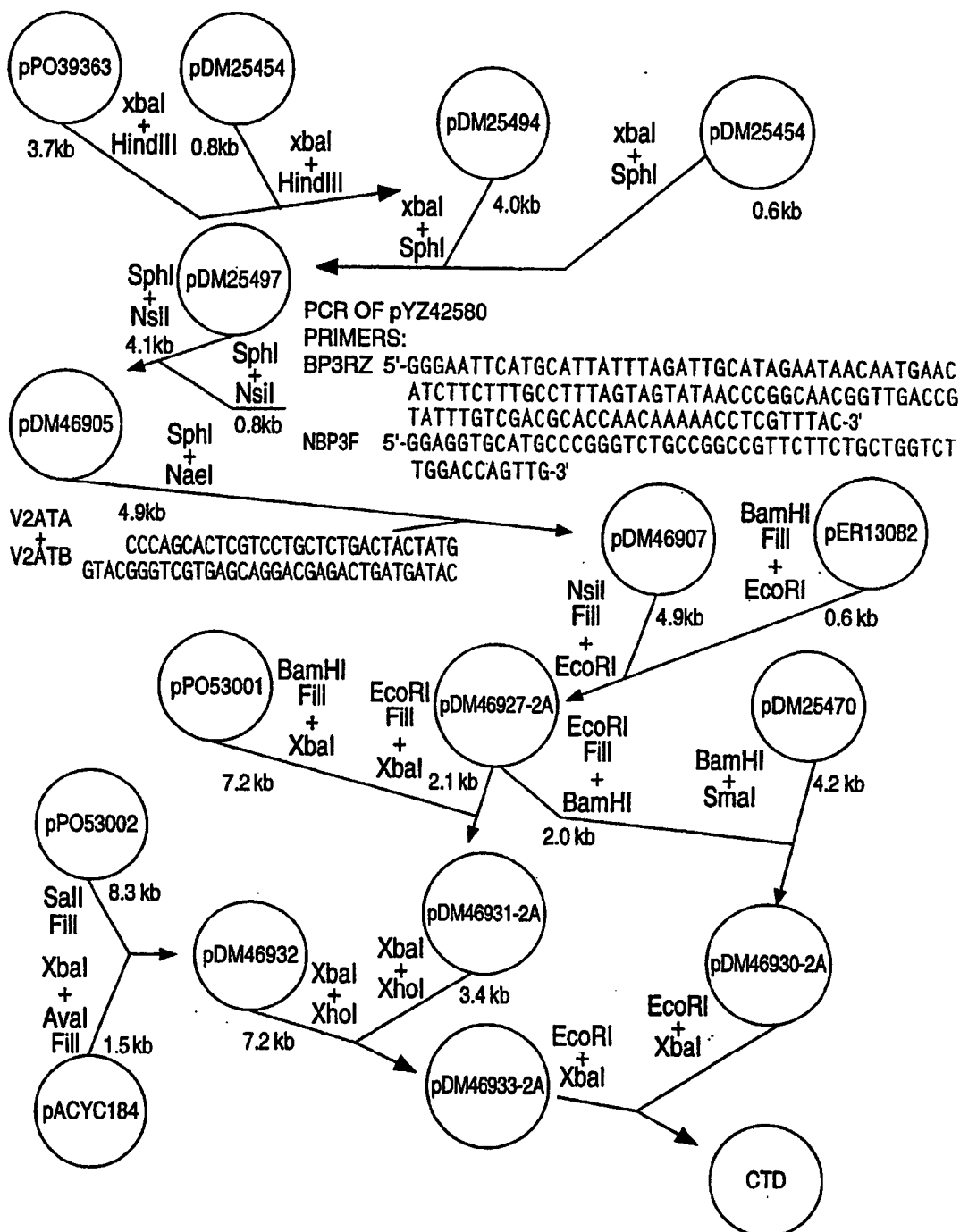
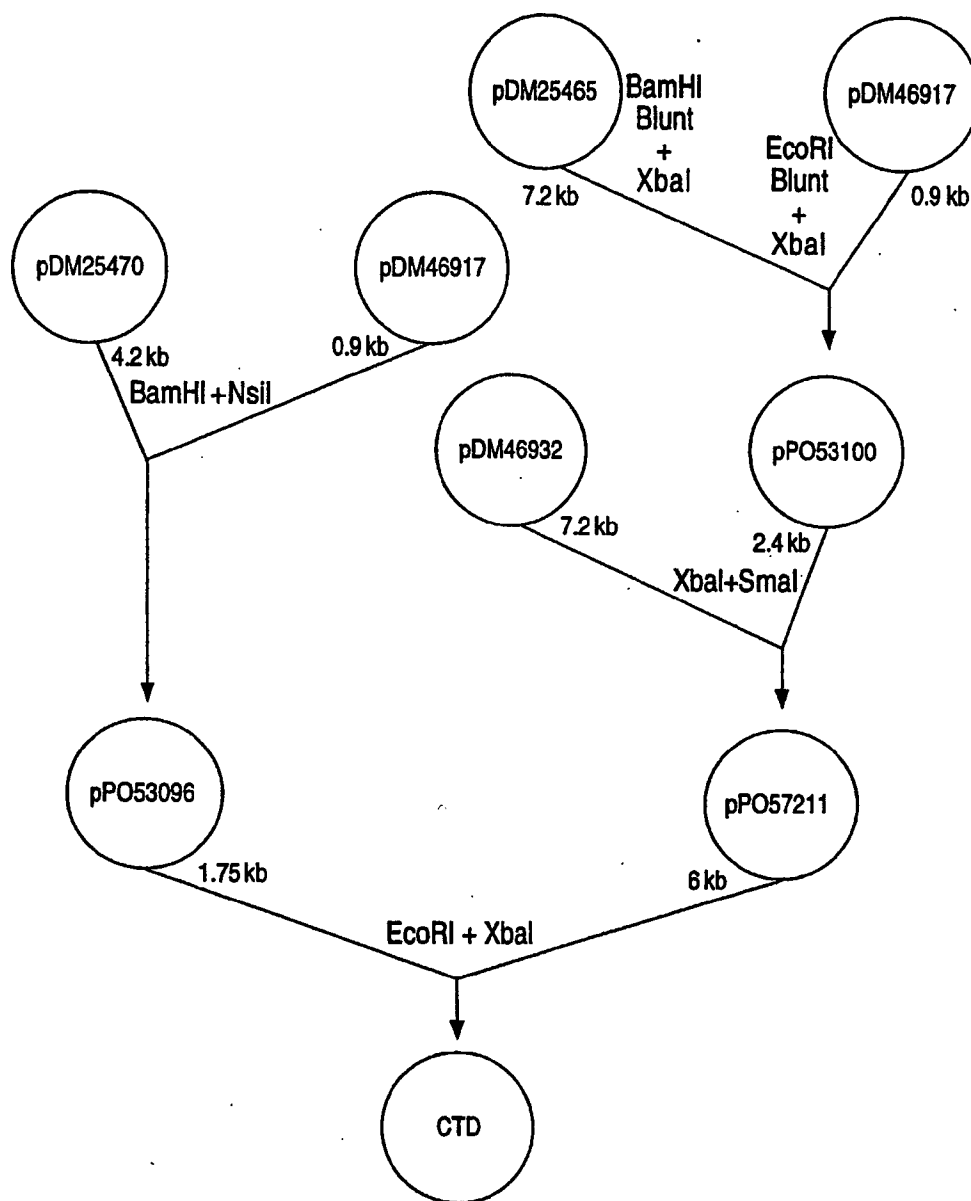


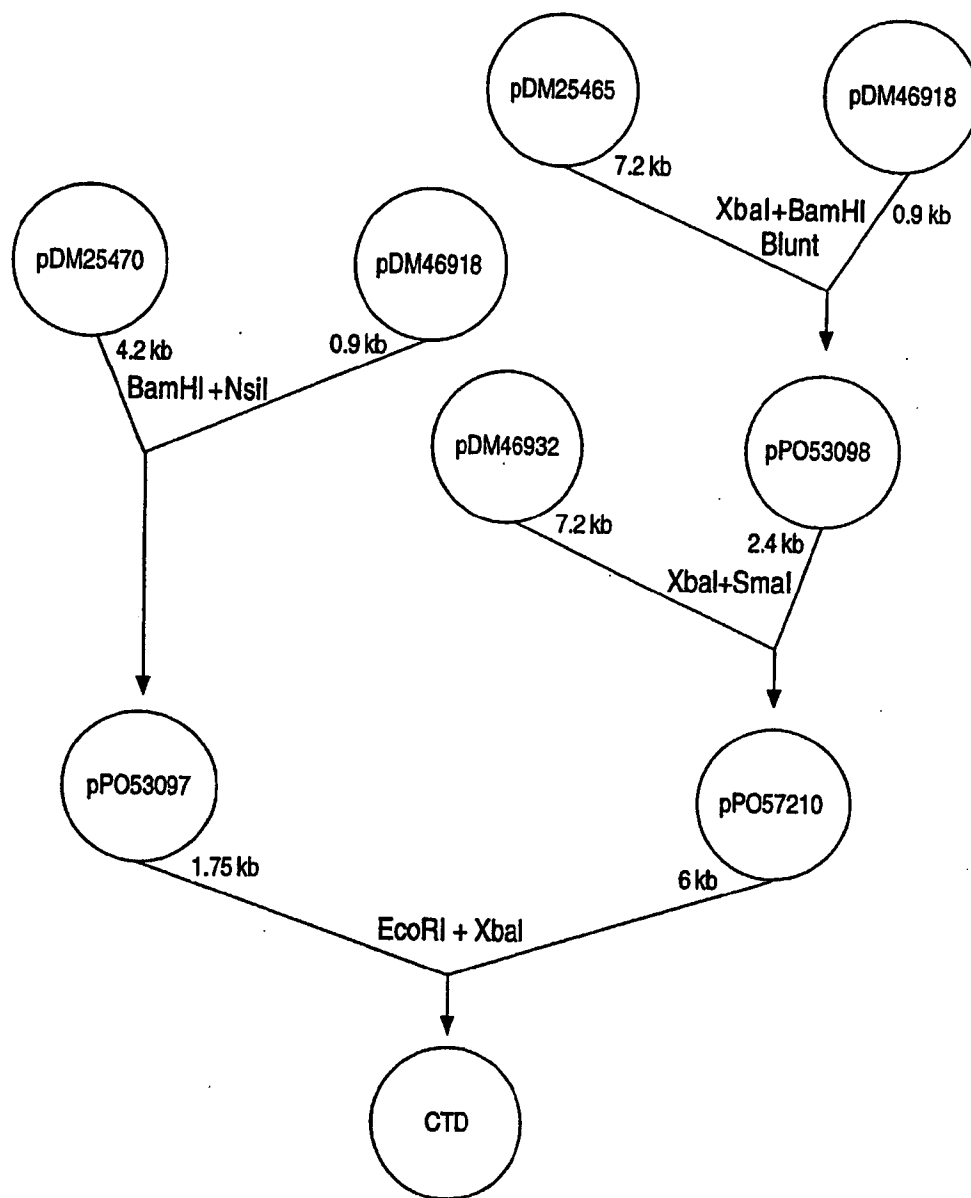
FIG. 18

SUBSTITUTE SHEET (RULE 26)

19/26

**FIG. 19**

20/26

**FIG. 20**

21/26

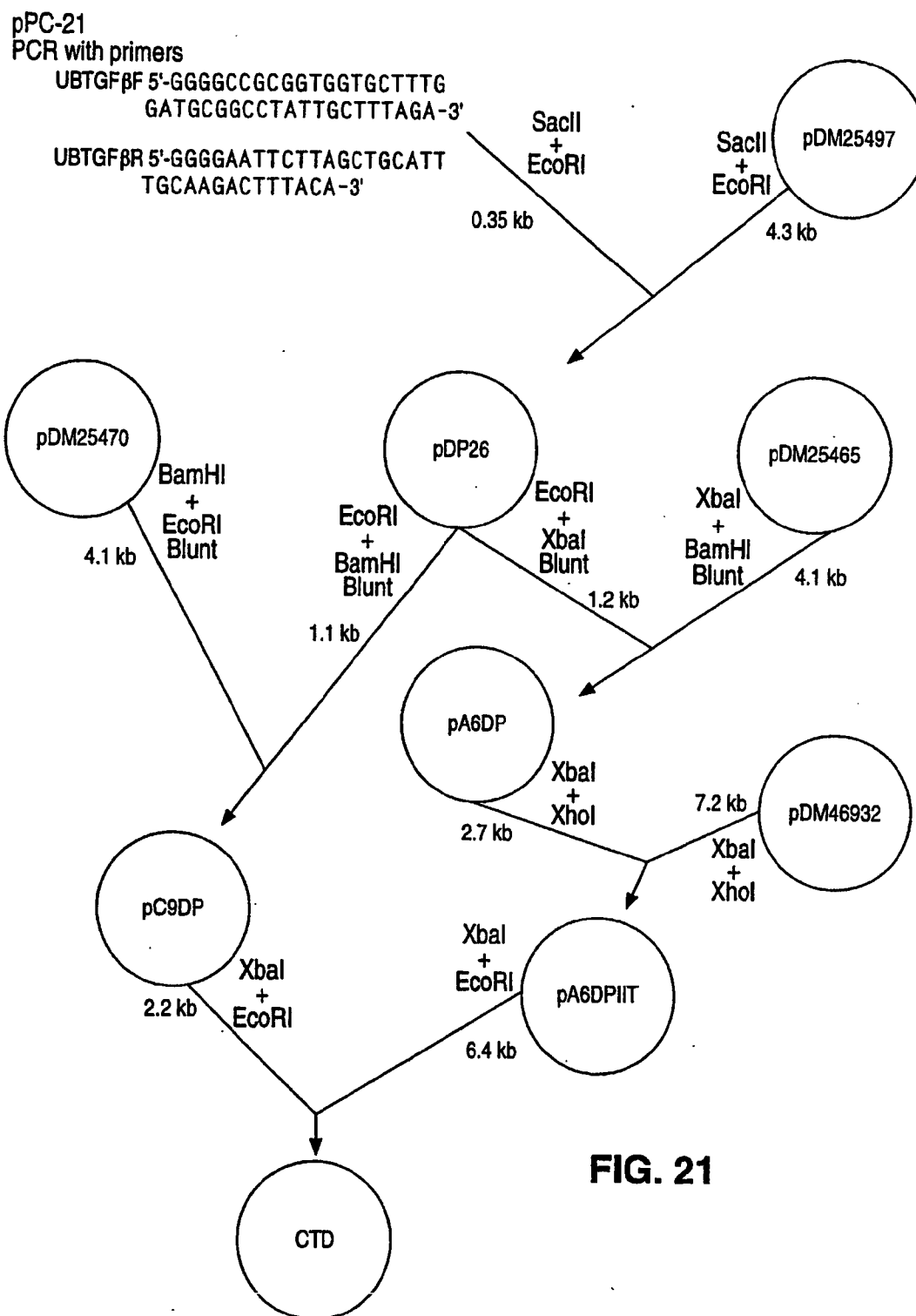


FIG. 21

22/26

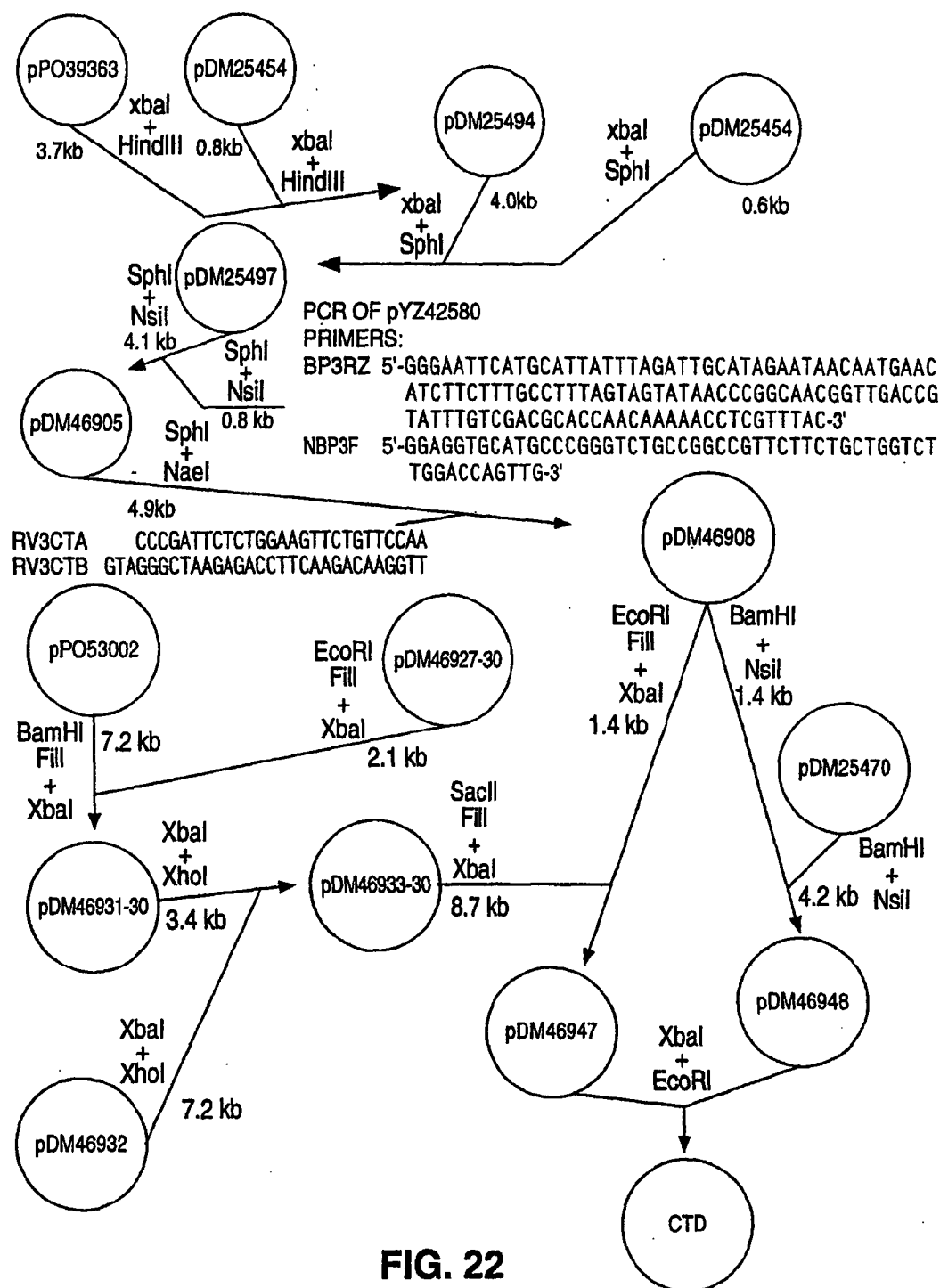


FIG. 22

23/26



FIG. 23

24/26

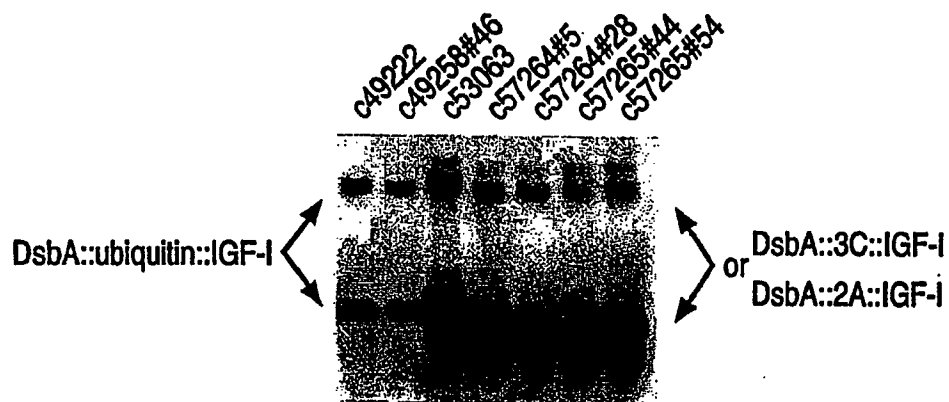


FIG. 24

25/26

FIG. 25A

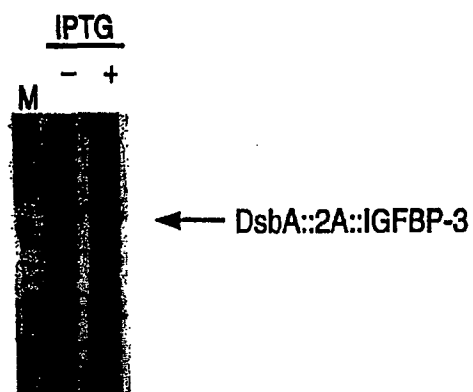
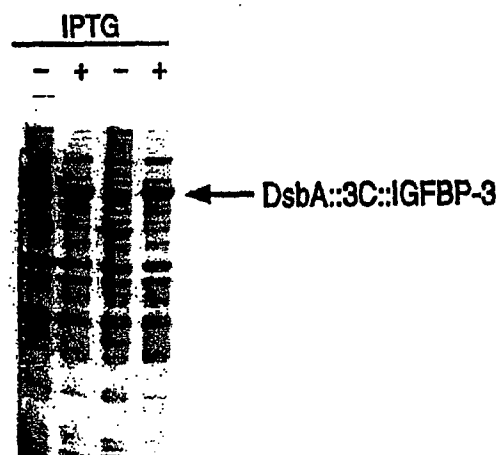


FIG. 25B



26/26

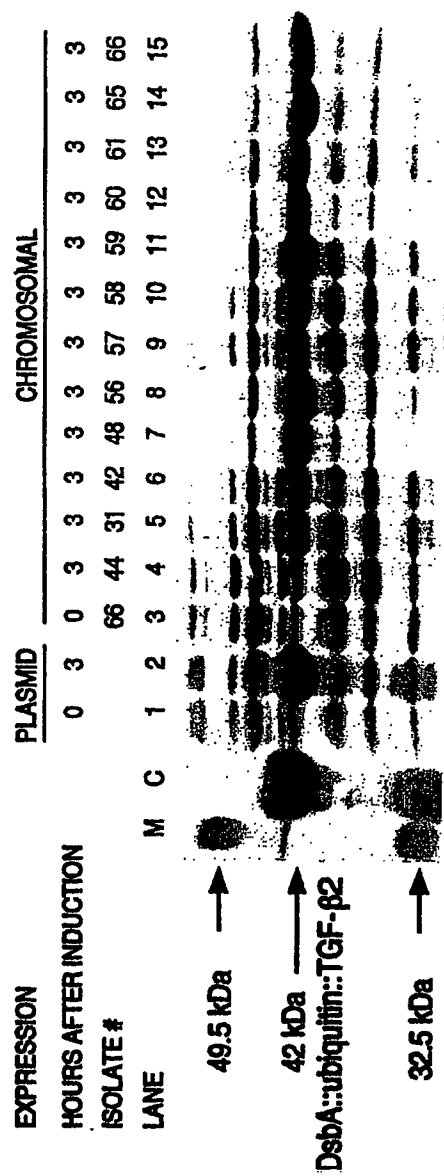


FIG. 26

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US96/09006

A. CLASSIFICATION OF SUBJECT MATTER IPC(6) : C07H 21/04; C12N 15/12, 15/63, 15/64, 15/70, 15/74, 15/90 US CL : Please See Extra Sheet. According to International Patent Classification (IPC) or to both national classification and IPC		
B. FIELDS SEARCHED Minimum documentation searched (classification system followed by classification symbols) U.S. : 435/69.1, 71.1, 172.1, 172.3, 252.3, 252.33, 320.1; 536/23.1, 24.1; 935/22, 27, 33, 38, 41, 42, 52, 72 Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched Electronic data base consulted during the international search (name of data base and, where practicable, search terms used) APS, BIOSIS, EMBASE, MEDLINE, DERWENT search terms: chromosom? transfer?, integrat?, express?, protein?		
C. DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	DIEDERICH et al. New Cloning Vectors for Integration into the λ Attachment Site <i>attB</i> of the <i>Escherichia coli</i> Chromosome. Plasmid. 1992, Vol. 28, pages 14-24, see entire document.	1-12
A, P	US 5,470,727 A (D. MASCARENHAS ET AL.) 28 November 1995 (28.11.95), see entire document.	1-12
<input type="checkbox"/> Further documents are listed in the continuation of Box C. <input type="checkbox"/> See patent family annex.		
* Special categories of cited documents:	"T"	Inter document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"A" document defining the general state of the art which is not considered to be of particular relevance	"X"	document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
"E" earlier document published on or after the international filing date	"Y"	document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	"&"	document member of the same patent family
"O" document referring to an oral disclosure, use, exhibition or other means		
"P" document published prior to the international filing date but later than the priority date claimed		
Date of the actual completion of the international search 24 JULY 1996	Date of mailing of the international search report 02 AUG 1996	
Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C. 20231 Facsimile No. (703) 305-3230	Authorized officer NANCY J. DEGEN Telephone No. (703) 308-0196	

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US96/09006

A. CLASSIFICATION OF SUBJECT MATTER:
US CL :

435/69.1, 71.1, 172.1, 172.3, 252.3, 252.33, 320.1; 536/23.1, 24.1; 935/22, 27, 33, 38, 41, 42, 52, 72